Supporting Information for Predicting Electronic Structure Properties of Transition Metal Complexes with Neural Networks

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Text S1: supporting files index

In addition to results reported here, additional files are provided as follows:

```
provided files:
  geometries.zip
     - csd_optimized_geos
      - test_[test number]_[% HFX]_[HS/LS].xyz
   - optimized_geos
      ID_[metal]_[oxidation]_[spin]_[ligands][% HFX].xyz
  - data.zip
   - ann_weights
       - split (splitting energy)
          [name]_[w/b (weight/bias)][layer number].csv
        - slope (HFX sensitivity)
          (same)
        - hs (high-spin bond length)
          (same)
       - 1s (low-spin bond length)
          (same)
- raw_data
     - split (splitting energy)
        — raw_data.csv<descriptors of all data>
        - all_results.csv <ANN pred. and data for all runs>
      - test_results.csv <ANN pred. and data for test runs>
     - slope (HFX sensitivity)
      (same)
     - hs (high-spin bond length)
      (same)
     - ls (low-spin bond length)
      (same)
     - csd_metrics (error and distance metrics for csd)
       - error_comp_split.csv
        - error_comp_ls.csv
       - error_comp_hs.csv
```

Text S2: Estimation of τ

We determine a representative value of τ by maximizing the log predictive likelihood of the corresponding GP based on the training data, which is a measure of how likely the observed data are under the GP, and is approximated [6] by

$$\log p\left(\mathbf{y}\left(\mathbf{x}_{n}\right)|\mathbf{x}_{n},\mathbf{X},\mathbf{Y}\right) \approx \log \left[\sum_{j=1}^{J} e^{-\frac{1}{2}\tau \left\|\tilde{\mathbf{y}}(\mathbf{x}_{n})-\tilde{\mathbf{y}}_{j}(\mathbf{x}_{n})\right\|_{2}^{2}}\right] - \log J - \frac{1}{2}\log 2\pi - \frac{1}{2}\log \tau^{-1}$$
(S1)

In the application here, we have only scalar output and we use the training data to optimize equation S1 with respect to τ numerically. We use J = 100 repeats, as in the network itself. The determined values of τ , based on the respective training data, are 0.4 for predicting the splitting energy, 0.07 for predicting the HF exchange sensitivity, and 10000 for the metal-ligand distances respectively. The magnitude of these numbers are close to the training errors observed, with $\sqrt{0.4^{-1}} \sim 2.5, 0.4^{-1} \sim 1.6, \sqrt{0.007^{-1}} \sim 12, 0.4^{-1} \sim 0.01$. These numbers represent the estimated inherent variance in the training data that limit of accuracy that could be expected from the trained networks.

Text S3: Use of Coulomb matrix descriptor

We compare the descriptors proposed in this work to the Coulomb matrix descriptor, which has previously [22, 9] been correlated with various molecular properties for a number of organic molecule data sets. In order to allow comparison of complexes with differing numbers of atoms, we pad all matrices with zeros to a size of of 151×151 , necessitating $\mathcal{O}(10^4)$ elements per compound. We sort the rows and columns of the matrices in order to obtain indexing-invariant representations and use KRR with an exponential kernel and the matrix L₁ norm as a distance metric as in Ref [32]. The complexes in our training data range in size from 7 to 151 atoms, but have a mean and median size of 38 and 29 atoms respectively. This large skew toward smaller complexes means that most of the descriptor elements are zero, and this may make learning good model parameters difficult. For example, the L_1 distances between the sorted matrix representations of the small Fe(III)(CN)₆ complex and two large complexes, Fe(III)(tbuc)₆ and Fe(III)(pisc)₆ are very similar (36.85 to 36.88 where the range of distances spans $\sim 20-60$), despite pisc being a similar strong C-conntecting ligand and tbuc being a much weaker Oconnecting ligand. We train and test on the same data as used in the other methods, but because the Coulomb matrix representation does not encode any functional-dependent information, we also provide a comparison against only B3LYP data (as opposed to varying HFX fractions).

Text S4: Testing ANN performance in molSimplify

In order to asses if the ANN can assist automated structure design, we used it to predict bond lengths instead of using the metal-ligand bond length databse integrated into our structure generation toolbox, molSimplify [13]. We selected four of the original benchmark structures where molSimplify was found not to reduce RMS gradient error relative to simple force fields. Further details about the test cases are in the original paper. We project the negative of the energy gradient on the metal and connection atoms at the initial geometry onto the vector joining them, as explained in Figure S32, and use this a measure of how close to an equilibrium bond length the initial geometry is. Note that a negative value for g means the bond would shrink in a steepest descent step, while a positive value means that it would lengthen. Large magnitudes indicate the bonds are far from equilibrium.

We achieve reductions in the absolute magnitude of g by 54-90% for bidentate cases and 7% for the monodentate case (Table S26). We note that the reductions in the metal-ligand projected gradient do not necessarily correspond to reductions in the RMS gradient, which considers contributions from all atoms. In the $Cr (bipy)_3$ case, the RMS gradient is reduced by 30%, but it is unchanged or marginally higher in the other cases. This may be explained by considering the signs of the projected gradient, which show that the ANN universally reduces the metal-ligand bond length relative to original structure. This brings the bidentate ligands closer to the metal center and hence closer to each other, and we observe that the dominant contribution to the RMS gradient is from other atoms in the ligand structure. This could possibly be improved by training a similar ANN on the bite angles.

Text S5: Molecular descriptors for CSD compounds

The poor correlation ($R^2 = 0.1$) between the Tanimoto dissimilarity (for CSD and training ligands) and the prediction error can be understood by considering that the molecular fingerprint is insensitive to the arrangement of groups in the ligand, so two ligands might appear similar in the Tanimoto metric because they both contain certain groups, but this does not ensure that the same groups are coordinating to the metal center. The descriptors used in this work strongly suggest that the immediate metal environment determines behavior of the complex, and so this highlights a specific difficulty in translating established ideas from organic molecular similarity analysis to transition metal systems.

Text S6: Dissimilarity metrics for LS/HS bond length prediction

Using the same dissimilarity metrics that were employed to evaluate reliability of spin-state splitting, correlations between HS bond distance error and proximity to test data is smaller for both the HS bond distances ($R^2 = 0.0, 0.1$ and 0.2 for the Tanimoto similarity metric, Pearson, and Euclidean distances, respectively) and LS bond distances ($R^2 = 0$ for all metrics). However, we do observe that four of the five large (i.e., > 0.1 Å) HS bond distance errors have a minimum Euclidean distance greater than 1.0, supporting the use of this heuristic for evaluating prediction reliability. Bond length errors are generally smaller for LS states compared to HS states, with only two cases (tests 26 and 30) greater than 0.1 Å. We observe an overall correlation between the low spin bond distance prediction inaccuracy and poor splitting energy prediction, but bond lengths may still be well-predicted when spin-state splittings are not (e.g., 0.006 - -0.03 Åerrors in LS bond distances for the cyclams).

Table S1: Ligand properties

Number	ID	Name	Denticity	Charge	Connection	$\max \delta \chi$	Bond Order	Truncated Kier
1	cl	chloride	1	1-	Cl	0	0	0
2	scn	thiocyanate	1	1-	S	0.03	2	2
3	pisc	t-butylphenyl isocyanide	1	0	С	-0.49	3	2.25
4	misc	Methyl isocyanide	1	0	С	-0.49	3	2
5	cn	cyanate	1	1-	С	-0.49	3	0
6	со	carbonyl	1	0	С	-0.89	3	0
7	ncs	isothiocyanate	1	1-	N	0.49	2	2
8	ammo	ammonia	1	0	Ν	0.84	1	0
9	bipy	2,2-bipyridine	2	0	Ν	0.49	2	4.297
10	phen	phenanthroline	2	0	Ν	0.49	2	3.868
11	en	ethylenediamine	2	0	Ν	0.84	1	3
12	porphyrin	porphyrin	4	2-	Ν	0.49	2	6.958
13	h2o	water	1	0	0	1.24	1	0
14	acac	acetylacetonate	2	0	0	0.89	2	3.10
15	tbuc	t-butyl catecholate	2	2-	0	0.89	1	2.52
16	OX	oxalate	2	2-	0	0.89	2	2.22
Number	ID	SMILES						
1	cl	[CI-]						
2	scn	[S-]C#N						
3	pisc	CC(C)(C)C1=CC=C(C=C	1)[N+]#[C-]					
4	misc	C[N+]#[C-]						
5	cn	[C-]#N						
6	со	CO						
7	ncs	[N-]=C=S						
9	bipy	C1ccnc(c1)c2ccccn2						
10	phen	C1=CC2=CC=C3C=CC=I	C1=CC2=ĆC=C3C=CC=NC3=C2N=C1					
11	en	NCCN						
12	porphyrin	[NH]1C2=CC3=NC(=CC4	=CC=C([NH	1]4)C=C5C:	=CC(=N5)C=C	1C=C2)C=	C3	
13	h2o	0		- /				
14	acac	CC(=O)CC(=O)C	CC(=O)CC(=O)C					
15	tbuc	CC(C)(C)C1=CC(=C([O-]))	CC(C)(C)C1=CC(=C([O-])C=C1)[O-]					
16	ox	[O-]C(=O)C([O-])=O	/					

Table S2: Core homoleptic ligands

motal	ovidation	number of converged & included HFX values per ligand										
metai	UXIUALIUIT	acac	bipy	c2h3s	cn	со	en	h2o	ncs	nh3	ох	total
CO	2	7	7	7	0	7	7	7	7	7	7	63
со	3	7	7	7	7	7	7	7	7	7	7	70
fe	2	7	0	7	7	7	7	7	7	7	7	63
fe	3	7	7	7	7	7	7	7	7	7	7	70
mn	2	7	7	7	0	7	7	7	7	7	5	61
mn	3	7	7	7	7	7	7	7	0	7	7	63
cr	2	7	7	7	0	7	7	7	0	7	7	56
cr	3	7	7	7	7	7	7	7	7	0	7	63
ni	2	7	7	7	5	7	7	7	7	7	7	68
total (core) 83 complexes, 577 HFX values, 1154 geometries												
total (a	additional)	11	1 com	plexes, 7	'68 H	FX v	alues	s, 1536	geon	netries		
total (all)		194 complexes, 1345 HFX values, 2690 geometries										

Table S3: List of test structures from the CSD, CSD IDs, metal and oxidation state along with a short note for each and references to the original observations

number	CSD ID	Metal	Ox.	Con. Atom (Ax./Equit)	Note	Reference
1	CODZAW10	Со	2	C/N	dicyano-cobyrinate	[19]
2	DORLEB	Co	2	C/N	tetrapyrrole	[7]
3	KOCLET	Cr	3	CI/O	tetrahydrofuran dichloride	[14]
4	KEHXEA	Co	3	CI/N	phthalocyanine	[25]
5	ZUNSEI	Co	3	CI/N	cylcam	[29]
6	ABZACO10	Co	2	O/O	benzoate	[5]
7	LUWPAU	Fe	2	N/N	phthalocyanine	[15]
8	TPYFEC04	Fe	2	CI/N	tetrapyridine	[36]
9	TPPSFE10	Fe	3	S/N	substitued porphyrin	[21]
10	FUMJOO	Mn	2	O/O	large oxygen ligand	[8]
11	BUHKIA	Ni	2	O/O	sulfoxide, diphenyl-propandionate	[18]
12	SUMLET	Cr	3	C/N	cyclam	[28]
13	BOSDIX	Fe	3	O/O	bidentate oxygen	[37]
14	BMADFE10	Fe	3	S/N	cyclam	[1]
15	AHAVUB	Mn	2	N/O	dinitrobenzoate	[16]
16	DOCNFE	Fe	2	C/N	benzyldioximate	[35]
17	AFAROO	Mn	3	C/N	cyclam	[23]
18	AHDNIC	Ni	2	O/N	cyclam	[3]
19	BUPTAH	Fe	3	O/O	bidentate oxygen	[31]
20	DEDKII	Fe	2	O/N	nitrogen rings and oxygen	[11]
21	PUTHIX	Fe	2	N/N	bidentate nitrogen	[39]
22	AGIZEX	Fe	2	N/N	bidentate nitrogen	[26]
23	AKAGEY	Fe	3	N/N	large monodentate nitrogen	[34]
24	GUWYUS	Fe	3	N/N	monodentate nitrogen ligand	[20]
25	EGILOW	Fe	3	O/N	heavily substituted porphyrin	[30]
26	BINPET	Cr	3	O/N	cyclam, oxygen	[2]
27	JUSCIL	Fe	2	N/N	bidentate nitrogen	[12]
28	BULVAG	Mn	2	O/N	salen-like	[4]
29	FIXGID	Mn	3	O/N	cyclam	[24]
30	KUSKEQ	Ni	2	O/O	monodentate oxygen	[33]
31	AGUWEE	Mn	2	O/N	cyclic nitrogen	[17]
32	DMAZCO03	Co	2	N/N	imidazole	[38]
33	YUJCIQ	Cr	3	C/N	cyclam	[27]
34	ACALEW	Cr	3	N/N	monodenate nitrogen	[10]
35	AKAGAU	Cr	3	N/N	monodentate nitrogen	[34]

number	spin	spin energy tolerance, au					
HFX = 0.20							
2	LS	$1.05 imes 10^{-5}$					
2	HS	$5 imes 10^{-2}$					
1	LS	$5 imes 10^{-6}$					

Table S4: Relaxed geometry optimization tolerances for some CSD structures .

Table S5: Excluded structures due to spin contamination

metal	ох	axlig	eqlig	aHF	metal	ох	axlig	eqlig	aHF
cr	3	nh3	nh3	all (7)	fe	2	tbuc	tbuc	0.30
mn	3	pisc	h2o	0.30	mn	2	ох	ох	0.25, 0.30
cr	3	tbuc	tbuc	0, 0.05					
total 5 complexes, 13 HFX values									

Table S6: Excluded structures due to geometric brekaup

_

metal	ох	axlig	eqlig	aHF	metal	ох	axlig	eqlig	aHF
CO	3	nh3	СО	0	ni	2	cn	cn	0,0.05
mn	3	ncs	ncs	all (7)	cr	2	ncs	ncs	all (7)
mn	2	cn	cn	all (7)	со	2	cn	cn	all (7)
cr	2	cn	cn	all (7)	cr	2	ncs	pisc	0
fe	3	со	scn	0,0.05	fe	2	ncs	pisc	0.25
ni	2	nh3	cn	all (7)	ni	2	scn	ох	all (7)
ni	2	h2o	h2o	all (7)					
total			13 com	13 complexes, 63 HFX values					

Table S7: List of input space descriptors and the normalization constants used in the ANNs. For a given variable *x*, the normalization is $\tilde{x} = \frac{x-c}{f}$.

unit	С	f
kcal/mol	-54.19	142.71
kcal/mol.HFX	-174.20	161.58
Å	1.8146	0.6910
Å	1.8882	0.6956
	2	1
	0	0.3
	-2	2
	-2	2
	1	1
	1	3
	-5.34	12.54
	-0.89	2.09
	0	3
	0.00	3
	0.00	4.29
	0.00	6.96
	unit kcal/mol kcal/mol.HFX Å Å	unit c kcal/mol -54.19 kcal/mol.HFX -174.20 Å 1.8146 Å 1.8882 2 0 -2 -2 -2 1 1 1 -5.34 -0.89 0 0 0.00 0.00 0.00 0.00

Table S8: Variable Selection for ΔE_{HS-LS} : set **a**. Values are given for regularized and unregularized coefficients and MSE in kcal²/mol²

	unregularized	\mathcal{L}_1 regularization
(Intercept)	7.13	1.14
metalcr	-23.3	-17.3
metalfe	-8.43	-2.76
metalmn	-25.4	-19
metalni	-29.8	-23.9
OX	6.73	6
alpha	-69.4	-64.4
axligbipy	1.82	3.48
axligc2h3ns	29.2	24.4
axligcl	-9.54	-4.77
axligcn	11.7	12.5
axligco	8.79	4.07
axligen	-4.65	0
axligh2o	-10.3	-9.9
axligncs	-10.2	-4.6
axlignh3	4.86	0.21
axligox	-22	-0.42
axligphen	-1.86	0
axligpisc	1.23	0
axligporphyrin	0	0
axligscn	-1.19	0
axligtbuc	-8.8	0
eqligbipy	9.07	0
eqligc2h3n	-1.19	1.37×10^{-5}
eqligcl	-6.67	-2.43
eqligcn	8.06	10.2
eqligco	11.1	12.7
eqligen	10.1	0.284
eqligh2o	-4.94	-4.31
eqligncs	-3.36	-0.298
eqlignh3	-0.946	0
eqligox	9.04	0
eqligphen	15.8	6.99
eqligpisc	34.2	31.2
eqligporphyrin	35.3	13.2
eqligscn	-4.83	0
eqligtbuc	-4.26	-1.58
axlig charge	-9.92×10 ⁻²	5.36
eqlig charge	-2.29	0
axlig dent	4.05	0
eqlig dent	-7.86	0
MSE	199	213

Table S9: Variable Selection for ΔE_{HS-LS} : set b .	Values are given for regularized and unregular-
ized linear coefficients and MSE in kcal ² /mol ²	

	unregularized	\mathcal{L}_1 regularization
(Intercept)	20.6	18.3
metalcr	-22.3	-17.9
metalfe	-7.03	-2.25
metalmn	-24	-19.1
metalni	-28.6	-23.6
OX	6.63	6.1
alpha	-69.5	-65
axlig charge	3.55	2
eqlig charge	-0.613	0.107
axlig dent	6.83	0.999
eqlig dent	-0.633	0
axlig connectCl	-12.9	-9.59
axlig connectN	-10.2	-6.14
axlig connectO	-17.1	-12.4
axlig connectS	-4.6	-2.39
eqlig connectCl	-23.1	-21.2
eqlig connectN	-18.2	-16.1
eqlig connectO	-23.6	-21.5
eqlig connectS	-21.6	-18.8
axlig natoms	-0.235	0
eqlig natoms	0.525	0.414
MSE	218	227

Table S10: Variable Selection for ΔE_{HS-LS} : set **c**. Values are given for regularized and unregularized linear coefficients and MSE in kcal²/mol²

	unregularized	\mathcal{L}_1 regularization
(Intercept)	26.4	13.6
metalcr	-22.5	-18.6
metalfe	-7.1	-3.18
metalmn	-24	-20
metalni	-28.9	-24.7
OX	6.6	6.17
alpha	-69.4	-66.1
axlig charge	3.57	2.34
eqlig charge	-1.22	0.702
axlig dent	8.6	0
eqlig dent	-0.841	0
axlig connectCl	-17.3	-7.43
axlig connectN	-19	-0.818
axlig connectO	-28.9	-4.73
axlig connectS	-9.04	0
eqlig connectCl	-26	-16.8
eqlig connectN	-24.6	-5.78
eqlig connectO	-31.5	-8.28
eqlig connectS	-24.6	-14
axlig natoms	-0.299	0
eqlig natoms	0.568	0.399
$\Sigma \Delta \chi$	-0.203	-1.95
$min\Delta\chi$	7.08	0
$max\Delta\chi$	5.43	-1.33
MSE	216	230

Table S11: Variable Selection for ΔE_{HS-LS} : set **d**. Values are given for regularized and unregularized linear coefficients and MSE in kcal²/mol²

	unregularized	\mathcal{L}_1 regularization
(Intercept)	26.1	14.8
metalcr	-22.5	-18.9
metalfe	-7.1	-3.49
metalmn	-24.1	-20.3
metalni	-28.8	-25.1
OX	6.58	6.2
alpha	-69.4	-66.4
axlig charge	3.69	2.39
eqlig charge	-1.3	0.595
axlig dent	7.92	0.115
eqlig dent	-0.48	0
axlig connectCl	-15.2	-7.78
axlig connectN	-15.2	-1.6
axlig connectO	-23.9	-5.75
axlig connectS	-7.03	0
eqlig connectCl	-27.3	-17.8
eqlig connectN	-26.9	-7.23
eqlig connectO	-34.5	-10.1
eqlig connectS	-25.8	-15.1
axlig natoms	-0.277	0
eqlig natoms	0.556	0.407
$max\Delta\chi$	-1.79	-1.47
$\Sigma \Delta \chi$	2	-1.68
MSE	216	228

Table S12: Variable selection for ΔE_{HS-LS} : set **e**. Values are given for regularized and unregularized linear coefficients and MSE in kcal²/mol²

	unregularized	\mathcal{L}_1 regularization
(Intercept)	32.7	19.1
metalcr	-22.2	-19.8
metalfe	-7.72	-4.74
metalmn	-24.1	-21.5
metalni	-29.6	-26.4
OX	6.6	6.38
alpha	-69.3	-67.3
axlig charge	5.12	2.83
eqlig charge	-3.68	0
axlig dent	6.49	1.89
eqlig dent	0.769	0.283
axlig connectCl	-16.7	-10.8
axlig connectN	-13.6	-4.97
axlig connectO	-18.1	-8.77
axlig connectS	0.368	0
eqlig connectCl	-26.5	-19.7
eqlig connectN	-30.7	-14
eqlig connectO	-46.1	-20.4
eqlig connectS	-37.8	-21.2
axlig natoms	-4.3×10^{-2}	0
eqlig natoms	0.32	0.378
axlig bo	3.64	1.39
eqlig bo	-5.96	-1.78
$\Sigma \ \Delta \chi$	2.25	-0.591
$max\Delta\chi$	-2.41	-1.46
MSE	213	221

Table S13: Variable selection for ΔE_{HS-LS} : set **f**. Values are given for regularized and unregularized linear coefficients and MSE in kcal²/mol²

	unregularized	\mathcal{L}_1 regularization
(Intercept)	27.8	17.8
metalcr	-22.4	-19.5
metalfe	-6.83	-3.71
metalmn	-24.3	-21.1
metalni	-29.9	-26
OX	6.69	6.27
alpha	-69.4	-66.9
axlig charge	2.38	2.11
eqlig charge	-0.265	0.992
axlig dent	11.3	2.18
eqlig dent	-2.95	0.846
axlig connectCl	-17.1	-9.14
axlig connectN	-14.4	-4.04
axlig connectO	-25.7	-9.6
axlig connectS	-1.94	-0.457
eqlig connectCl	-28.6	-21.5
eqlig connectN	-33.9	-14.6
eqlig connectO	-39.4	-17.7
eqlig connectS	-35.4	-22
axlig ki	-2.65	-0.343
eqlig ki	4	1.78
$max\Delta\chi$	-1.19	-0.721
$\Sigma \Delta \chi$	2.48	-0.868
MSE	218	227

Table S14: Variable selection for ΔE_{HS-LS} : set **g**. Values are given for regularized and unregularized linear coefficients and MSE in kcal²/mol²

	unregularized	\mathcal{L}_1 regularization
(Intercept)	35.2	21.6
metalcr	-22.2	-19.6
metalfe	-7.5	-4.28
metalmn	-24.4	-21.5
metalni	-30.1	-26.5
OX	6.63	6.32
alpha	-69.3	-67.1
axlig charge	4.04	2.75
eqlig charge	-2.84	0
axlig dent	10.1	2.19
eqlig dent	0.186	2.29
axlig connectCl	-17.7	-10.4
axlig connectN	-13.9	-4.73
axlig connectO	-22	-8.81
axlig connectS	0.214	0
eqlig connectCl	-27.7	-21.3
eqlig connectN	-34	-16.3
eqlig connectO	-47.9	-23.7
eqlig connectS	-41.9	-25.8
axlig bo	2.05	1.24
eqlig bo	-5.52	-2.71
axlig ki	-1.66	0
eqlig ki	1.97	0.925
$max\Delta\chi$	-2.08	-1.09
$\Sigma \Delta \chi$	2.41	-0.622
MSE	215	223

Table S15: Variable selection for $\frac{\partial \Delta E_{H-L}}{\partial a_{HF}}$: set **a**. Values are given for regularized and unregularized coefficients and MSE in HF kcal²/mol²HFX²

	unregularized	\mathcal{L}_1 regularization
(Intercept)	-167	-167
metalcr	21.7	21.2
metalfe	-26.3	-26.3
metalmn	-15.2	-15.2
metalni	14.9	14.2
OX	28.2	27.9
eqligbipy	-20.7	-16.6
eqligc2h3ns	-30.5	-30.1
eqligcl	-2.95	-0.898
eqligcn	-6.33	-4.87
eqligco	-18.4	-14.8
eqligen	-4.02	-1.31
eqligh2o	8.22	11.8
eqligncs	1.03	2.08
eqlignh3	4.88	7.82
eqligox	-0.272	-1.45
eqligphen	-1.28	0
eqligpisc	-31.6	-27.5
eqligporphyrin	8.4	1.34
eqligscn	6.26	6.82
eqligtbuc	10.8	7.53
axligbipy	15.5	10.2
axligc2h3ns	-1.39	-0.326
axligcl	6.15	0.329
axligcn	-9.2	-14
axligco	-21.8	-24.1
axligen	17.4	13.5
axligh2o	15	11.8
axligncs	7.92	2.13
axlignh3	4.47	2.07
axligox	11	0.491
axligphen	-10.2	-11.8
axligpisc	-16.3	-18.9
axligporphyrin	0	0
axligscn	2.97	-1.65
axligtbuc	8.32	0
eqlig charge	2.31	0
axlig charge	-1.12	-4.04
eqlig dent	-1.75	0
axlig dent	-8.15×10 ⁻²	0
alpha	5.72	5.42
MSE	422	423

Table S16: Variable selection for $\frac{\partial \Delta E_{H-L}}{\partial a_{HF}}$: set **b**. Values are given for regularized and unregularized linear coefficients and MSE in HF kcal²/mol²HFX²

	unregularized	\mathcal{L}_1 regularization
(Intercept)	-189	-186
metalcr	21.9	21.7
metalfe	-26.9	-26.7
metalmn	-16.3	-16.1
metalni	13.8	13.3
OX	28.6	28.2
eqlig charge	-0.266	-0.359
axlig charge	-3.79	-3.49
eqlig dent	0.179	0
axlig dent	-7.42	-6.37
eqlig natoms	-0.47	-0.452
axlig natoms	2.13×10^{-2}	0
eqlig connectCl	10.5	9.61
eqlig connectN	17.6	17
eqlig connectO	19.3	18.9
eqlig connectS	20.1	19.2
axlig connectCl	18.6	18.1
axlig connectN	24.5	23.8
axlig connectO	29.7	28.8
axlig connectS	18.1	17.7
alpha	4.89	4.58
MSE	472	472

Table S17: Variable selection for $\frac{\partial \Delta EH-L}{\partial a_{HF}}$: set **c**. Values are given for regularized and unregularized linear coefficients and MSE in HF kcal²/mol²HFX²

	unregularized	\mathcal{L}_1 regularization
(Intercept)	-170	-167
(Intercept)	0	0
metalcr	20.8	20.9
metalfe	-27.6	-27.2
metalmn	-17.1	-16.8
metalni	13.1	12.8
OX	28.6	28.4
eqlig charge	-2.06	-1.99
axlig charge	-3.98	-3.44
eqlig dent	2.38	0
axlig dent	0.717	0
eqlig natoms	-0.385	-0.325
axlig natoms	-0.144	-0.161
eqlig connectCl	-2.56	0.619
eqlig connectN	-9.37	-2.65
eqlig connectO	-13.9	-5.88
eqlig connectS	6.76	9.66
axlig connectCl	6.83	7.52
axlig connectN	-0.514	2.22
axlig connectO	-4.21	0
axlig connectS	5.29	6.36
alpha	4.21	3.69
$\Sigma\Delta\chi$	1.76	1.02
$max\Delta\chi$	14.4	13.1
min $\Delta\chi$	14.2	13
MSE	449	450

Table S18: Variable selection for $\frac{\partial \Delta E_{H-L}}{\partial a_{HF}}$: set **d**. Values are given for regularized and unregularized linear coefficients and MSE in HF kcal²/mol²HFX²

	unregularized	\mathcal{L}_1 regularization
(Intercept)	-168	-166
metalcr	20.8	20.9
metalfe	-27.4	-27.2
metalmn	-17.2	-17
metalni	13.2	12.8
OX	28.6	28.4
eqlig charge	-2.24	-2.05
axlig charge	-3.61	-3.24
eqlig dent	0.574	-0.291
axlig dent	-1.65	-1.14
eqlig natoms	-0.388	-0.347
axlig natoms	-0.108	-0.147
eqlig connectCl	-4.31	0
eqlig connectN	-12.4	-6.75
eqlig connectO	-18.8	-11.6
eqlig connectS	5.36	8.2
axlig connectCl	10.9	10.6
axlig connectN	7.03	8.26
axlig connectO	5.61	7.47
axlig connectS	9.34	9.41
alpha	4.17	3.55
$\Sigma\Delta\chi$	5.86	4.97
$max\Delta\chi$	0.542	0.725
MSE	451	452

Table S19: Variable selection for $\frac{\partial \Delta E_{H-L}}{\partial a_{HF}}$: set **e**. Values are given for regularized and unregularized linear coefficients and MSE in HF kcal²/mol²HFX²

	unregularized	\mathcal{L}_1 regularization
(Intercept)	-190	-187
metalcr	20.7	20.6
metalfe	-26.5	-26.4
metalmn	-16.9	-16.8
metalni	13.9	13.5
OX	28.6	28.3
eqlig charge	0.243	0
axlig charge	-4.29	-3.67
eqlig dent	0.362	-1.06
axlig dent	-3.11	-1.84
eqlig natoms	-0.252	-0.172
axlig natoms	-6.34×10^{-2}	-0.159
eqlig connectCl	-4.69	-0.473
eqlig connectN	-9.57	-3.89
eqlig connectO	-8.03	-0.743
eqlig connectS	16.6	19.5
axlig connectCl	12.8	12.3
axlig connectN	8.55	9.11
axlig connectO	7.62	8.08
axlig connectS	10.1	9.26
alpha	5.12	4.43
axlig bo	5×10^{-1}	0
eqlig bo	5.57	5.77
$\Sigma \Delta \chi$	5.62	4.81
$max\Delta\chi$	1.4	1.45
MSE	444	445

Table S20: Variable selection for $\frac{\partial \Delta E_{H-L}}{\partial a_{HF}}$: set **f**. Values are given for regularized and unregularized linear coefficients and MSE in HF kcal²/mol²HFX²

	unregularized	\mathcal{L}_1 regularization
(Intercept)	-192	-122
metalcr	20.9	17.9
metalfe	-27.7	-21.6
metalmn	-16.8	-10.8
metalni	14.3	1.98
OX	28.5	19.8
eqlig charge	-1.58	-0.317
axlig charge	-3.9	-2.2
eqlig dent	9.74	0
axlig dent	-0.587	0
eqlig connectCl	-1.05	0
eqlig connectN	-0.537	0
eqlig connectO	-5.67	0
eqlig connectS	16.9	0
axlig connectCl	15.5	0
axlig connectN	13.8	0
axlig connectO	15.2	0
axlig connectS	11.4	0
n alpha	5.18	0
axlig ki	-0.177	0
eqlig ki	-4.11	0
$max\Delta\chi$	0.434	0
$\Sigma\Delta\chi$	3.91	4.27
MSE	453	553

Table S21: Variable selection for $\frac{\partial \Delta E_{H-L}}{\partial a_{HF}}$: set **g**. Values are given for regularized and unregularized linear coefficients and MSE in HF kcal²/mol²HFX²

	unregularized	\mathcal{L}_1 regularization
(Intercept)	-192	-187
metalcr	20.9	20.8
metalfe	-27.7	-27.5
metalmn	-16.8	-16.6
metalni	14.3	13.8
OX	28.5	28.2
eqlig charge	-1.58	-1.68
axlig charge	-3.9	-3.66
eqlig dent	9.74	7.56
axlig dent	-0.587	0
eqlig connectCl	-1.05	0
eqlig connectN	-0.537	0
eqlig connectO	-5.67	-3.96
eqlig connectS	16.9	15.7
axlig connectCl	15.5	14.3
axlig connectN	13.8	13.3
axlig connectO	15.2	14.4
axlig connectS	11.4	11.3
alpha	5.18	4.64
axlig ki	-0.177	-0.364
eqlig ki	-4.11	-3.37
$max\Delta\chi$	0.434	0.522
$\Sigma\Delta\chi$	3.91	3.83
MSE	453	454



Figure S1: Comparison of errors for different descriptor sets for a regularized linear effects model predicting ΔE_{H-L} in kcal/mol (left) and $\frac{\partial \Delta E_{H-L}}{\partial a_{HF}}$ in kcal/molHFX (right). Set **a** includes the metal properties and full ligand identity and number of atoms. Set **b** replaces ligand identity with the identity of connection atom only, while set c adds information from the sum, maximum and minimum ligand $\delta \chi$ to set **b**. Set d is the same as set c but excludes the minimum $\delta \chi$. Set **e** adds in bond order information with an MSE, while set f replaces the ligand size metric with our truncated index. Set **g** represents our final set, and includes the same descriptors from **f** and adds bond order information.

Table S22: Optimal hyperparameter selection for KRR and SVR models found via a grid search and 10-fold cross-validation.Paremteres were selected usnig a cartesian grid search in $[10^{-7}, 10^4]$ for the regularization weights, $[10^{-4}, 10^3]$ for the exponential kernel correlation length and [0.1, 0.9] for ν .

	σ	λ/C (KRR/SVR)	ν
	(kernel lengthscale)	(regularization weight)	(SV fraction)
KRR (set g)	1	10^{-4}	
SVR (set g)	1	100	0.75
KRR (sorted Coulomb Matrix, B3LYP)			
KRR (sorted Coulomb Matrix, B3LYP)	316	0.01	

Figure S2: Binary ground state classification tree for heteroleptic compounds. M indicates metal identity, I ligand connection atom, o oxidation state, a the fraction of HF exchange and me the sum of $\delta\chi$ values across ligands. The first line in each leaf node is the percent of elements in that leaf that have the indicated ground state, and the second line indicates the percentage of the total heteroleptic population in each leaf node. Dashed blue arrows indicate yes, solid red arrows indicate no.



Key:	
% HS % of pop.	% LS % of pop.



Figure S3: Model predictions of ΔE_{H-L} and data for Mn using an ANN. The ligands are described by two numbers indicating the equatorial first and then the axial, color coded by ligand identity (green for halogen, gray for carbon, blue for nitrogen, and red for oxygen). The error bars represent an estimated 1 standard deviation from the mean prediction.



Figure S4: Model predictions of $\Delta E_{\text{HS-LS}}$ and data for Co using an ANN. The ligands are described by two numbers indicating the equatorial first and then the axial, color coded by ligand identity (green for halogen, gray for carbon, blue for nitrogen, and red for oxygen). The error bars represent an estimated 1 standard deviation from the mean prediction.



Figure S5: Model predictions of ΔE_{H-L} and data for Cr using an ANN. The ligands are described by two numbers indicating the equatorial first and then the axial, color coded by ligand identity (green for halogen, gray for carbon, blue for nitrogen, and red for oxygen). The error bars represent an estimated 1 standard deviation from the mean prediction.



Figure S6: Model predictions of ΔE_{H-L} and data for Fe(II) (top) and Ni(II) (bottom) using an ANN. The ligands are described by two numbers indicating the equatorial first and then the axial, color coded by ligand identity (green for halogen, gray for carbon, blue for nitrogen, and red for oxygen). The error bars represent an estimated 1 standard deviation from the mean prediction.

Table S23: Splitting energy predictions and data in kcal/mol, compared for the full set of complexes. Each complex is named [metal oxidation axial ligand eq equatorial ligand]. For each complex, the first row has the data obtained from DFT using varying fractions of exact exchange $(a_{\rm HF})$ and the second row has the ANN predictions

		0.00	0.05	a	HF	0.00	0.05	0.00
cr 2 eq h2o ax ncs	data	-12.6	-14.5	-15.7	-16.7	-17.7	-18.6	-19.5
	pred	-8.37	-13.1	-15.4	-16.9	-18.4	-19.9	-21.8
cr 2 eq porphyrin ax nh3	data	13.8	11.9	9.98	8.13	5.99	5.55	5.32
cr 2 eq scn ax nisc	data	11.9 5.14	0.567	-4	8.58	6.9 -12 7	-16.9	-20.8
	pred	7.04	2.99	-2.5	-7.76	-11.8	-15.1	-18.3
cr 2 eq ncs ax pisc	data	11.3	8.27	3.43	-1.33	-5.82	-10	-13.9
cr 2 eq cn av nh3	data	8.4 5.23	6.51 -1.62	3.61	-0.0798	-3.86	-6.58	-8.45
	pred	4.81	0.785	-2.55	-5.35	-7.93	-10.2	-12
cr 2 eq tbuc ax tbuc	data	7.45	4.37	1.46	-1.26	-3.83	-6.25	-8.56
cr 2 eq phen av sch	data	4.1	3.19	1.94	-0.0318	-3.06	-6.91	-10.9
or 2 og prior ax son	pred	8.11	4.4	0.0103	-4.36	-8.82	-13.5	-18.4
cr 2 eq en ax co	data	9.12	4.19	-0.386	-4.62	-8.51	-12.1	-15.3
cr 2 eq biny ax biny	pred	10.5	3.83	-1.79	-5.36	-7.91	-11.2	-13.9
	pred	9.12	3.51	-0.407	-2.82	-4.71	-6.48	-8.2
cr 2 eq en ax en	data	-7.2	-8.91	-10.5	-12	-13.4	-14.7	-16
or 2 og og og og	pred	-3.58	-6.9	-9.09	-10.8	-12.2	-13.2	-13.9
	pred	36.4	34.4	31.7	27.9	23.2	18.4	14.0
cr 2 eq pisc ax pisc	data	33.2	28	23.2	18.7	14.4	10.5	6.85
	pred	29.7	26.7	23.1	18.8	15	12	9.41
01 2 EY UX AX 1120	pred	-29.5	-32.5	-32,8	-34.9	-41.4	-44.5	-38.4
cr 2 eq h2o ax cl	data	-26.6	-27.4	-27.6	-28.3	-29.6	-30.2	-30.7
or 0 og ph2 ov oop	pred	-23.8	-25.2	-26.4	-27.5	-28.5	-29.4	-30.3
ci 2 eq illis ax scri	pred	-12	-14	-13.7	-16.7	-10.0	-20.1	-21.4
cr 2 eq ox ax ox	data	8.53	5.44	2.32	-0.786	-3.89	-6.92	-9.9
or 2 og b2o ov b2o	pred	4.47	2.89	0.835	-1.65	-4.57	-7.76	-10.8
CI 2 eq 1120 ax 1120	pred	-14.5	-9.85	-13.5	-13	-19.1	-20	-21
cr 2 eq cn ax h2o	data	19.2	18.5	17.8	16.9	15.9	14.9	13.8
or 2 og pige ev pee	pred	16.6	16.4	16.3	16	15.5	14.5	11.8
ci 2 eq pisc ax rics	pred	-	30.5	25.5	20.1	14	7.89	0.0285
cr 2 eq acac ax acac	data	3.66	-0.746	-4.83	-8.78	-15.9	-18.4	-20.4
or 2 og biny ov b2o	pred	1.43	0.00453	-2.61	-6.79	-10.8	-14.2	-17.6
Ci 2 eq bipy ax 1120	pred	-6.42	-6.85	-12.3	-9.69	-17.2	-14.8	-21.1
cr 2 eq nh3 ax nh3	data	-10.2	-11.6	-13	-14.3	-17	-18.1	-17.8
	pred	-6.81	-9.52	-11.4	-12.8	-14.3	-16.2	-19.1
	pred	22.0	17	13.2	9.44	4.00 6.42	3.68	0.934
cr 2 eq c2h3ns ax c2h3ns	data	31	26.5	22.2	18.1	14.1	10.4	6.8
or 2 og phon ov phon	pred	31.3	28.4	24.5	19.9	14.9	9.81	4.93
ci 2 eq phen ax phen	pred	5.99	0.882	-2.44	-4.63	-2.55	-5.65 -8.34	-0.32
cr 2 eq tbuc ax nh3	data	-11.8	-14.1	-16.5	-18.7	-21	-23.1	-25.2
or 2 og ph2 av op	pred	-11.2	-12.5	-13./	-15.2	-1/.4	-20.1	-22.6
	pred	0.612	-2.48	-5.21	-7.28	-8.86	-10.9	-14
cr 3 eq c2h3ns ax c2h3ns	data	-18.7	-20	-21.2	-22.3	-23.4	-24.3	-25.3
cr 3 eq acac ay acac	data	-15.5	-1/./	-19.1	-20.3	-21.5	-22.5	-23.1
	pred	-10.1	-14.4	-16.8	-17.8	-18.5	-19.2	-20
cr 3 eq h2o ax scn	data	-19.2	-20.5	-21.6	-22.8	-23.9	-25.5	-26.5
cr 3 eq en ax nh3	data	-17.3	-20.2	-21.7	-22.7	-23.5	-24 -27 1	-24.4
	pred	-20.4	-21.7	-22.4	-22.9	-23.3	-23.7	-24.2
cr 3 eq scn ax scn	data	-21.4	-22.8	-24	-25.2	-26.2	-27.2	-28.1
cr 3 eq en ax en	data	-17.3	-19.3	-20.3	-21.3	-22.6	-24.4	-26.4
	pred	-22.2	-22.5	-22.7	-22.9	-23	-23.2	-23.4
cr 3 eq cn ax cn	data	-19.5	-20.7	-21.8	-22.8	-23.8	-24.7	-25.6
cr 3 eq cn av nes	data	-13.8	-15.1	-16.3	-17.4	-18.9	-20.5	-22
010 04 011 07 1100	pred	-13.6	-17	-19.8	-21.8	-23.7	-26.3	-28.5
cr 3 eq tbuc ax tbuc	data	-	-	-21.9	-22.8	-23.7	-24.6	-25.4
cr 3 eq nos av nos	pred	-17.2	-18.6	-21.3	-21.8	-22.1	-22.6	-23.4
or o eq nos ax nos	pred	-14.6	-15.5	-16.5	-17.6	-18.9	-20.2	-21.4
cr 3 eq ox ax ox	data	-20.7	-21.6	-22.5	-23.3	-24.2	-24.9	-25.7
cr 3 eq cl ax cl	pred	-18	-19.2	-20.3	-21.2	-21./	-22	-22.5
	pred	-12.7	-14	-15.8	-18.5	-21.4	-23.4	-24.7
cr 3 eq co ax co	data	-21.5	-22.6	-23.5	-24.4	-25.3	-26.1	-26.9

	tuno	0.00	0.05	0.1	a _{HF}	0.20	0.25	0.20
name	pred	-16.7	-18.6	-19.6	-20.4	-21	-21.5	-21.9
cr 3 eq h2o ax co	data	-18.2	-19.6	-20.8	-22	-23.1	-24.1	-25.1
cr 3 eg bipy ax bipy	data	-15.6 -21.4	-16.8	-17.7	-18.5	-19.2	-19.8	-20.3
-	pred	-15.9	-18.7	-20.2	-21	-21.5	-21.9	-22.2
cr 3 eq ncs ax cl	data	-18.8	-20.5	-21.6	-22.7	-23.7	-24.7	-25.7
cr 3 eq tbuc ax pisc	data	-5.9	-9.35	-12.8	-15.9	-15.1	-20.3	-21.9
	pred	-8.48	-10.2	-12	-13.7	-15.2	-16.6	-17.9
ci s eq ox ax ci	pred	-20.4	-21.3	-22.2	-23	-23.0	-24.5	-23.5
cr 3 eq h2o ax h2o	data	-21.2	-22.1	-23.1	-24	-24.8	-25.7	-26.5
cr 3 eg ox ax pisc	data	-12.4	-12.1	-19.4	-20.1	-20.8	-19.8	-22.1
	pred	-7.49	-10.1	-12.1	-13.8	-15.6	-17	-18.2
mn 2 eq acac ax nn3	pred	-4.7 -5.43	-12.8	-20.1	-26.6	-31.7	-36.3	-40.3
mn 2 eq nh3 ax nh3	data	-21.4	-25.4	-28.9	-32.6	-36	-39.3	-42.5
mn 2 eg co ax co	data	-16.6	-21.8	-27.2	1.68	-35.7 -6.57	-38.9	-40.9 -21.4
	pred	24	18.8	11.8	2.64	-5.38	-10.2	-14.2
mn 2 eq cn ax nn3	pred	-3.47	-3	-5.6 -6.64	-11.7	-17.4	-22.7	-27.6
mn 2 eq en ax en	data	-15.7	-20.1	-24.3	-28.3	-32.1	-38.2	-39
mn 2 eg acac ax acac	pred data	-15.5 -10.1	-20.1 -18.6	-23.9 -25	-27.2	-30	-32.7	-35.3 -43.9
	pred	-12.7	-16.9	-21.5	-26.4	-31.1	-34.8	-37.8
mn 2 eq en ax cn	data	3.26	-2.16 1.06	-7.31	-12.2 -13	-16.8 -19.7	-22.7	-28.6
mn 2 eq bipy ax h2o	data	-14.6	-20.3	-25.5	-30.2	-34.5	-38.5	-42.1
mn 2 eg h2o av h2o	pred data	-15.6	-21.3	-26.9 -42 1	-31.7	-35.2	-37.4	-39.1
	pred	-30.8	-34.3	-37.4	-40.4	-43.1	-45	-46.2
mn 2 eq ncs ax nh3	data	-12.4	-18	-23.1	-25.4	-29.9	-33.9	-37.8
mn 2 eq bipy ax bipy	data	4.7	-2.95	-9.9	-16.2	-21.9	-27	-31.6
	pred	0.523	-3.75	-9.54	-15.7	-20.3	-24.3	-28.1
	pred	-35.4	-41.7	-45.4	-47.3	-48.5		
mn 2 eq c2h3ns ax c2h3ns	data	43.7	34.5	25.6	17.2	9.17	1.6	-5.51
mn 2 eq ncs ax co	data	27.7	18.7	9.92	1.59	-6.26	-13.6	-20.5
	pred	21.9	17.4	9.61	1.72	-5.24	-12.3	-16.9
min 2 eq nes ax nes	pred	-23.9	-32	-35.1	-42.4	-40.8	-44.5	-47.1
mn 2 eq phen ax phen	data	3.97	-3.6	-10.5	-16.7	-22.3	-27.3	-31.8
mn 2 eg nh3 ax co	data	15.5	7.04	-1.01	-8.61	-15.7	-22.4	-28.5
	pred	12.8	6.28	0.325	-7.14	-14.5	-20	-23.8
min 2 eq en ax n20	pred	-23.8 -20.4	-27.4	-30.8	-34.∠ -35	-37.3	-40.3 -39.7	-43.2 -41.4
mn 2 eq en ax co	data	24.8	15.9	7.43	-0.592	-8.16	-15.3	-21.9
mn 2 eg nh3 ax pisc	data	4.37	-2.39	-8.82	-1.33	-8.26	-14.6	-20.8
	pred	2.92	-2.32	-8.51	-15.5	-23.2	-28.4	-32.7
mn 3 eq en ax co	pred	2.03	-3.86	-6.09	-9.5 -8.05	-14.1	-15.5	-18.2
mn 3 eq phen ax cl	data	1.51	-1.15	-3.57	-5.85	-8.03	-10.6	-12.2
mn 3 eg phen ax phen	pred data	-3.49 10	-4.15 7.49	-5.02 5.17	-6.44	-8.3	-9.99	-11.3
	pred	7.47	5.32	3.61	1.77	-0.2	-2.13	-3.99
mn 3 eq cn ax nh3	pred	5.43	3.11 3.6	0.654	-1.68	-3.95 -3.27	-6.12	-8.21 -5.58
mn 3 eq pisc ax pisc	data	41.1	35.9	30.8	26	21.7	17.7	14.1
mn 3 eg ncs ax cl	data	37.4 -1.42	34.9 -4.14	-6.69	25.7 -9.08	-11.4	21.5 -13.5	19.4 -15.6
	pred	-3.51	-4.94	-6.59	-8.58	-10.9	-13	-14.6
mn 3 eq h2o ax co	data	2.61	0.268	-2.03	-4.21 -3.66	-6.34 -5.99	-8.39	-10.4 -9.77
mn 3 eq ox ax ox	data	-3.68	-5.71	-7.44	-9.07	-10.6	-12.2	-13.7
mn 3 eg co ax co	pred data	-3.51 17 1	-4.11 13.7	-5.14 10.4	-7.28	-10.2 4.55	-13.4 1.84	-17.1
	pred	17.3	14.8	12.1	9.32	7.06	5.26	3.76
mn 3 eq acac ax acac	data	2.92	0.269	-2.31	-4.3 -2.81	-6.33	-8.23	-10.1
mn 3 eq nh3 ax nh3	data	2.3	0.504	-1.46	-3.21	-4.95	-6.66	-8.38
mn 3 eg nh3 av co	pred	2.89	1.01	-0.523	-2.06	-3.95	-6.09	-8.09
	pred	4.27	1.18	-1.99	-4.59	-6.65	-8.49	-10.1
mn 3 eq cn ax cn	data	22	18.5	15.2	12.1	9.18	9.18	6.52
mn 3 eq nh3 ax scn	data	-16.5	-18.1	-19.6	-21.3	-23.1	-24.9	-26.7
mn 3 eg biny av biny	pred	-14.4	-17	-19.6	-21.7	-23.6	-25.9	-28.9
	pred	10	6.76	4.65	2.89	1.13	-0.754	-2.82
mn 3 eq h2o ax pisc	data	-8	-14.7	-21.2	-31.5	-36.5	-41	-
mn 3 eq en ax en	data	3.7	1.92	0.149	-23.2	-3.42	-20.0	-6.75
	pred	3.89	1.54	-1.02	-2.83	-4.02	-4.94	-5.86

	4	0.00	0.05	a	HF 15	0.00	0.05	0.00
name mn 3 eq phen ax nh3	data	-1.37	-3.2	-5.05	<u> </u>	-8.72	-10.5	-12.3
mn 3 eq h2o ax h2o	pred	1.99	-1.46	-3.93	-5.6	-6.81	-7.76	-8.61
	data	-5.06	-6.61	-8.16	-9.71	-11.3	-12.8	-14.3
mn 3 eq c2h3ns ax c2h3ns	pred	-4.55	-5.59	-6.52	-7.69	-9.32	-11.4	-13.7
	data	28.9	25.3	21.9	18.7	15.7	12.8	10.1
fe 2 eq co ax co	pred	26.7	22.2	19.4	16.8	14	10.8	7.61
	data	66.7	57.1	47.8	38.8	30.3	22.2	14.5
fe 2 eq acac ax acac	pred	60.4	53.8	45.4	37.5	29	21.2	15.4
	data	8.06	2.18	-3.18	-8.1	-20	-23.5	-23.7
fe 2 eq tbuc ax co	pred	3.11	-0.278	-4.38	-9.21	-14.3	-19.4	-24.9
	data	14.6	10.3	5.78	1.88	-3.22	-8.37	-4.03
fe 2 eq cn ax cn	pred	16.7	11	5.37	0.76	-3.93	-8.83	-13.9
	data	36.6	27.5	18.8	10.5	2.92	-4.09	-10.5
fe 2 eq tbuc ax tbuc	pred	36.7	26.4	16.7	7.32	0.181	-2.82	-4.33
	data	-17.6	-22.2	-26	-29.5	-32.6	-35.6	-
fe 2 eq h2o ax ncs	data	-18.9 -9.37	-24.4 -12.1	-27.8 -14.7	-29.5	-30.4 -19.9	-30.9 -22.5	-25
fe 2 eq en ax en	data	-13.8	-15.4 6.41	-16.6	-18.4 -1.39	-21.4 -5.03	-24.7	-27.5
fe 2 eq cl ax pisc	data	9.48 15.9	3.75 10.6	-0.159 5.14	-3.37 -0.496	-7.38 -6.23	-10.3 -13.4	-13.3 -18.8
fe 2 eq ncs ax h2o	pred data	16.7 - <u>6.22</u>	12.7 -9.95	8.18 - <u>13.6</u>	-17.6	-4.36	-10.8 -23.9	-15 -26.8
fe 2 eq pisc ax cl	data	-7.85 47.2	-11.6 40.6	-15.7 33.8	-20.1 27	-23.3 20.3	-26.2 13.8	-29 7.51
fe 2 eq phen ax phen	pred	44.6	36.1	30.1	24.6	17.8	12	7.84
	data	27.5	21	15.1	9.58	1.73	-2.67	-7.04
fe 2 eq scn ax h2o	pred	18.4	13.9	9.73	5.14	0.668	-3.22	-7.89
	data	-5.53	-8.55	-11.6	-14.8	-17.9	-17.3	-20.5
fe 2 eg nh3 ax nh3	pred	-6.56	-8.09	-9.57	-11.2	-13.3	-16	-18.6
	data	3.93	0.0674	-3.64	-7.18	-10.6	-13.8	-16.5
fe 2 eq c2h3ns ax c2h3ns	pred	2.23	-2.84	-6.52	-10	-12.8	-14.5	-15.9
	data	75.1	66.6	58.2	50.2	42.2	34.4	27.2
fe 2 eq ox ax pisc	pred	69.8	64.8	57.8	49.3	40.1	32.8	26.3
	data	28.8	22.9	17	11.3	5.53	-0.106	-8.43
fe 2 eq pisc ax cn	pred	22.4	19.4	15	8.85	1.78	-4.02	-9.1
	data	81.2	73.2	65.8	57.4	57.4	41	33.2
fe 2 eq acac ax cn	pred	69.1	67	64.7	60.3	51	38.9	30.4
	data	16.2	9.91	4.21	-1.67	-4.58	-11.6	-14.9
fe 2 eq ox ax ox	pred	12.8	7.53	1.76	-3.31	-8.46	-13.4	-16.2
	data	-13.7	-18.6	-22.7	-26.4	-29.8	-33	-35.9
fe 2 eg h2o ax h2o	pred	-18.5	-21.9	-25.2	-27.4	-28.6	-29.2	-29.7
	data	-14	-16.9	-20.1	-23.1	-26.1	-28.9	-31.5
fe 2 eq ncs ax pisc	pred	-11.4	-16.5	-21.1	-23.8	-25.7	-27.4	-29.1
	data	20.3	15	9.12	3.37	-1.55	-	-4.39
fe 2 eg en ax co	pred data	14.4 41.2	10.6 33.5	6.23 25.9	1.36 18.6	-2.84 11.6	5.18	-9.89 -0.805
fe 2 eq pisc ax ncs	pred	35.4	29.8	23	16.7	11.8	6.1	-0.0024
	data	47.6	33.4	27.6	21.5	17.4	11.3	5.4
fe 2 eq ncs ax co	pred	33.2	29.3	23.3	17.5	12.1	6.57	2.24
	data	28.3	23.2	17.6	11.8	5.85	0.102	-5.36
fe 2 eq ncs ax ncs	pred	29.9	22.1	14.8	10.1	6.06	1.72	-2.7
	data	-1.72	-6.89	-11.7	-16.2	-20.3	-24.2	-27.7
fe 3 eq tbuc ax tbuc	pred	-5.55	-9.9	-14.1	-18	-21.9	-25.7	-28.3
	data	-7.95	-11.1	-14.1	-17.2	-20.2	-23.2	-26.3
fe 3 eq cn ax cn	pred	-7.56	-10.7	-13.3	-15.9	-18.4	-20.7	-22.9
	data	57.9	51.8	46	40.3	34.8	29.4	24.2
fe 3 eq acac ax acac	pred	54.6	49.7	45.7	41.7	36.3	28.9	19.5
	data	8.94	5.47	2.02	-1.45	-4.97	-8.55	-12.2
fe 3 eq porphyrin ax h2o	pred	6.45	3.35	0.96	-0.856	-2.91	-5.33	-7.55
	data	18.7	15.3	12.1	8.76	-1.43	3.11	0.871
fe 3 eq h2o ax cn	pred	19.3	14.8	9.89	4.73	0.907	-1.44	-3.15
	data	14.7	11.6	8.5	5.36	2.1	2.78	-0.841
fe 3 eq porphyrin ax cn	pred	13.5	9.08	6.51	4.84	3.29	1.46	-0.922
	data	44.8	39.8	35	31.7	27	22.4	17
fe 3 eq co ax co	pred	38.8	36.8	34	30.1	25.5	20.6	16.2
	data	43.4	37.2	31.2	25.5	20	14.7	9.5
fe 3 eq ox ax ox	pred	45.5	36.8	30.4	24.7	19.3	15.1	12.2
	data	2.6	-0.574	-3.78	-7.03	-10.3	-13.7	-17.1
fe 3 eq pisc ax h2o	pred	2.68	-0.701	-4.73	-8.16	-11.4	-14.6	-17.8
	data	28.9	24.6	17.1	15	9.63	2.92	-1.87
fe 3 eq acac ax cl	pred	30.3	24.9	19.3	14.2	8.24	3.5	0.317
	data	1.11	-7.09	-10.2	-16	-16.2	-15	-18.2
fe 3 eq ox ax ncs	pred	-6.49	-7.3	-8.33	-9.96	-12	-14.2	-16.3
	data	8.73	4.93	1.26	-2.36	-5.81	-9.44	-13.1
fe 3 eg en ax en	pred	3.56	2.05	0.0278	-2.94	-6.08	-9.38	-13.4
	data	22.7	19.6	16.3	10.2	9.6	6.1	2.56
fe 3 eq pisc ax pisc	pred	18.8	15.4	12.2	9.55	6.96	4.6	2.45
	data	65.5	61.4	54.9	49.5	40.8	34.4	26.1
fe 3 eq bipy ax co	pred	59.1	56.7	52.9	46.3	38.7	31.9	27.3
	data	29.7	23.9	16.7	10.1	3.55	-2.55	-16.9
fe 3 eg pisc ax ncs	pred	29.9	24.7	19	11.9	4.95	-2.21	-8.63
	data	41.5	42.5	32.3	26.5	27.2	20.4	13.9
fe 3 eg ncs ax cl	pred	38.4	32.2	27.8	24.8	22.1	18.2	12.9
	data	1.61	-1.64	-10.5	-13.5	-16.5	-19.5	-22.7
fe 3 eg phen ax phen	pred	-0.111	-3.99	-8.3	-12.5	-15.8	-18.3	-20.4
	data	28.6	25.2	21.7	14.7	11	7.17	3.21
	pred	24.8	22.9	19.7	15.5	11.3	7.49	4.24

	tura 0	0.00	0.05	aj		0.00	0.05	0.00
fe 3 eg h2o ax h2o	data	-15.6	-18	-21	-23.4	-26	-28.7	-31.6
for Querral and a	pred	-9.04	-13.4	-17	-19.1	-20.6	-22.3	-24.4
te 3 eq cl ax cl	data pred	-13.9 -9.3	-16.6 -13.4	-26.3	-22.4 -21.9	-25.5 -26.6	-28.7 -30.2	-38.6 -32.4
fe 3 eq pisc ax cl	data	39.1	34.1	35.7	30.7	25.8	21.1	16.5
fe 3 eq c2h3ns ax c2h3ns	data	38	34.8 54.7	31.8 48.9	29.2 43.2	26.4	22.3 32.3	17.4 27
	pred	62.1	55.4	47.4	41.9	36.9	31.7	27.6
fe 3 eq nh3 ax scn	data	20.7	17.2	13.9	10.7	7.65	4.53	1.13
fe 3 eq acac ax ncs	data	-6.13	-9.36	-12.4	-15.4	-18.3	-21.1	-23.9
fe 3 eg co av sch	pred	-5.44	-9.91	-12.7	-14.3	-15.4	-16.1	-16.7
	pred			-10.9	-15.3	-18.3	-21	-23.7
fe 3 eq bipy ax bipy	data	29.7 25.3	29.4 23.9	25.3	21.1	16.9	12.7 9.78	8.53 5.89
fe 3 eq ncs ax cn	data	19.9	16	12.2	8.44	4.74	1.02	-2.73
fe 3 eg nos av nos	pred	20.2	15.9	12.1	8.41	4.56	1.19	-1.98
le 5 eq lics ax lics	pred	-2.02	-6.27	-9.38	-11.6	-13.2	-14.5	-15.9
fe 3 eq nh3 ax nh3	data	21	17.2	13.4	9.59	5.8	2.03	-1.73
fe 3 eq cl ax pisc	data	20.2	15.1	10.3	5.6	4.97	0.215	-4.42
	pred	19.1	15.6	10.9	5.82	2.28	0.0921	-1.74
	pred	-13.1	-14.5	-16.3	-18.3	-20.2	-20.5	-23.4
co 2 eq en ax en	data	3.16	0.284	-2.48	-5.19	-7.81	-10.4	-12.7
co 2 eq tbuc ax tbuc	data	-11.5	-13.7	-15.6	-17.5	-19.2	-20.8	-22.4
co 2 eg cl av nisc	pred	-13.8	-15.4	-17.1	-18.8	-20.4	-22	-23.5
co z eq ci ax pisc	pred	11	8.74	6.3	3.46	0.241	-3.54	-7.66
co 2 eq cn ax h2o	data	-30.3	-32.2	-34.6	-37.3	-40.1	-43	-46
co 2 eq en ax pisc	data	16.4	12.5	8.23	4.11	0.192	-3.49	-6.93
co 2 eg phen av h2o	pred	8.45	7.32	5.99	4.04	0.65	-3.56	-6.03
	pred	3.97	1.33	-1.55	-4.37	-6.51	-7.96	-9.16
co 2 eq phen ax phen	data	8.28	4.41	0.79	-2.61	-5.77	-8.74	-11.5
co 2 eq h2o ax ncs	data	-4.76	-6.41	-8.51	-10.2	-11.9	-13.5	-15
co 2 og c2h3ne av c2h3ne	pred	-7.4	-8.68	-9.82	-10.9	-12	-13.6	-16.5
	pred	32.7	25.4	19	13.6	9.01	4.8	1.71
co 2 eq nh3 ax nh3	data	0.576	-2.17	-4.44 -4.75	-7.45	-9.97 -11 7	-12.3	-14.4
co 2 eq acac ax cn	data	12.5	8.82	5.29	1.95	-0.987	-3.12	-6.87
co 2 eq nh3 ax h2o	pred data	11.1 7.28	7.41 4.48	3.23	-0.215 -1 14	-2.97	-5.27	-/.41 -9.11
	pred	1.72	-0.547	-2.54	-4.16	-5.73	-7.47	-9.44
co 2 eq acac ax acac	data pred	-3.04 -7.52	-6.6 -10	-9.73	-12.5	-14.9 -14.2	-13.2 -15.1	-15.4 -16
co 2 eq ncs ax ncs	data	-8.26	-10.8	-13.2	-15.4	-17.4	-19.2	-20.9
co 2 eg bipy ax bipy	data	-11.3 8.05	-14.7	-17	-18.2	-19.2	-20.5	-22.6
	pred	7.28	2.9	-0.524	-3.46	-6.07	-8.66	-11.9
co 2 eq co ax co	pred	28.1	22.3 19.6	16.6	9.27	6.23 4.86	1.48	-5.44 -1.13
co 2 eq en ax h2o	data	12.8	9.86	6.97	4.13	1.33	-1.41	-4.11
co 2 eg h2o ax h2o	data	-11.8	-14.2	-16.2	-18.1	-4.29	-0.41	-22.6
	pred	-10.9	-12.9	-15.3	-18.2	-21.4	-23.8	-25.3
	pred	-	52.4	49.3	47.5	40.9	36.2	32.0
co 3 eq acac ax nh3	data	29.4	30.2	27.7	25.2	22.7	20.2	17.8
co 3 eq tbuc ax pisc	data	11.8	9.19	6.23	-0.016	-3.36	-6.88	-11.7
co 3 eq nh3 av nh3	pred	10.2	8.16	5.98	2.95	-2.19	-9.67	-16.3
	pred	46.9	42.9	39.9	37.1	33.9	30.6	26.6
co 3 eq ox ax ox	data	-4.67	-7.87	-11.2	-14.6	-15.7	-21.6	-26
co 3 eq en ax en	data	49.6	47.7	45.4	42.8	40.1	37.3	34.4
00 3 00 2020 2X 2020	pred	46.7	43.9	40	35.4	31.2	28	24.9
co s eq acac ax acac	pred	29.3	29.5	25.4	23.1	20.4	16.7	10.5
co 3 eq phen ax phen	data	43.2 48	40.5	36.5	32.6 39	28.7	24.7	20.9
co 3 eq cn ax cn	data	88.5	83.6	78.7	73.8	69.1	64.5	60
co 3 eq c2h3ne av c2h3ne	pred	74.6	74	72.7	70.2	65.9 73.7	59.8 69.1	51.1
CO O EQ CENOIS AX CENOIS	pred	74.8	74.6	74.2	72.8	69.4	63.6	56.7
co 3 eq co ax co	data	77 71.8	71.6	66.3 64	61.2	56.3 54 5	51.6 49 9	47 45 1
co 3 eq tbuc ax tbuc	data	2.61	1.38	0.259	-0.943	-2.55	-4.74	-7.18
co 3 ea h2o av h2o	pred	5.13	3.38	1.48	-1.26	-5.64	-10.7	-15.6
60 3 EY 1120 at 1120	pred	11.5	8.36	5.78	3.39	1.51	0.0404	-1.79
co 3 eq ncs ax ncs	data	8.87	8.94	8.84	8.02	6.47	4.09	2.08

				ä	а _{нғ}			
name	type	0.00	0.05	0.1	0.15	0.20	0.25	0.30
	pred	15.6	12.6	9.55	5.42	1.15	-1.67	-4.04
co 3 eq pisc ax pisc	data	47.2	46.9	43.3	39.1	34.6	30.1	25.6
••	pred	46.2	44.3	42.2	39.4	35.9	32.1	28.2
co 3 eq bipy ax bipy	data	54.4	51.8	48.7	44.7	40.6	36.6	32.7
	pred	48.2	45.8	42.8	39.2	35.5	32.4	29.8
co 3 eq acac ax ncs	data	6.48	4.32	2.33	-0.259	-3.87	-7.95	-12.2
	pred	7.31	4.81	2.25	-0.448	-3.55	-6.93	-10.2
co 3 eq ox ax ncs	data	20.5	19.2	19.4	14.8	10.2	5.84	1.61
•	pred	21	18.3	15	11.5	8.37	5.4	2.52
co 3 eq acac ax co	data	23.7	19.6	15.8	11.5	7.16	2.7	-2.3
•	pred	18.4	16.5	14.1	11.6	9.14	6.06	2.15
ni 2 eg tbuc ax tbuc	data	-19.1	-22.3	-25.4	-28.5	-31.5	-34.4	-37.3
•	pred	-18.2	-21.8	-26.8	-30.8	-33.2	-34.9	-36.3
ni 2 eq bipy ax nh3	data	-13.2	-16.4	-19.2	-22	-24.9	-27.8	-30.7
	pred	-15.4	-18.3	-21	-23.7	-26.7	-29.6	-32
ni 2 eg en ax en	data	-18.8	-21.6	-24.3	-27.2	-30	-32.8	-35.6
•	pred	-24.7	-25.2	-25.7	-26.6	-28.3	-30.6	-33
ni 2 eq co ax co	data	-21	-24.4	-27.8	-31.3	-34.6	-37.9	-41.1
•	pred	-19.4	-22.8	-25.3	-28	-31	-34.3	-37.4
ni 2 eg nh3 ax nh3	data	0.552	-3.22	-6.79	-10.7	-14.5	-18.2	-21
•	pred	-4.75	-6.97	-10	-13.4	-16.1	-18.5	-21.3
ni 2 eq acac ax cn	data	-10.3	-13.4	-16.5	-19.5	-22.5	-25.4	-28.2
•	pred	-9.03	-12.4	-16.8	-21.2	-23.9	-25.2	-26.4
ni 2 eq pisc ax pisc	data	2.09	-1.92	-6.39	-10.6	-14.4	-18.2	-21.9
	pred	-2.05	-4.82	-7.77	-10.8	-13.9	-16.8	-19.8
ni 2 eg en ax h2o	data	1.41	-1.64	-4.68	-7.78	-10.8	-13.9	-17
•	pred	3.16	-1.42	-6.98	-11.8	-14.7	-16.6	-18.7
ni 2 eq bipy ax bipy	data	-17.1	-20.3	-23.5	-26.6	-29.8	-32.8	-35.8
	pred	-23.4	-24.3	-25	-25.8	-27.1	-28.9	-31.2
ni 2 eq ncs ax ncs	data	-32	-34.9	-37.8	-40.5	-43.1	-45.6	-47.9
	pred	-30.9	-33.4	-36	-39.6	-42.4	-43.9	-44.6
ni 2 eq cn ax cn	data	-	-	-20.4	-23.8	-27.1	-30.4	-33.5
•	pred			-20.5	-25.5	-28.8	-31.3	-33.8
ni 2 eq acac ax co	data	-13.3	-15.4	-17.8	-20.5	-23.3	-26.1	-28.8
	pred	-12.6	-14.9	-17.3	-20	-22.5	-24.7	-26.7
ni 2 eq c2h3ns ax c2h3ns	data	-19.5	-22.5	-25.6	-28.8	-32	-35.1	-38.1
	pred	-18.1	-22.1	-25.4	-28.1	-30	-32	-34.6
ni 2 eq acac ax nh3	data	-23.5	-25.9	-28.4	-31	-33.5	-36.1	-38.5
	pred	-22.4	-26.5	-29.5	-31.9	-33.8	-35.4	-36.7
ni 2 eq acac ax acac	data	-23.3	-26.8	-25.2	-28.9	-32.5	-35.9	-39.2
	pred	-23.4	-25	-26.9	-29.3	-32.3	-35	-37.2
ni 2 eq ox ax ox	data	-17.5	-21.2	-24.8	-28.6	-31.9	-35.1	-38.2
	pred	-17.9	-20.6	-24	-28.2	-31.8	-33.9	-35.3



Figure S7: Parity plot for 2 standard deviation from the mean prediction and absolute prediction error for test case ΔE_{H-L} prediction using ANN. All units are kcal/mol. The black line is y=x.



Figure S8: Normalized error histogram for HF = 0.2 (B3LYP) test data (top) and CSD structures (bottom), comparing ANN, KRR and SVR models using descriptor set \mathbf{g} , as well as a KRR model using the Coulomb matrix descriptor (trained on B3LYP data only.)

Table S24: RMS prediction errors for ΔE_{H-L} on test data using a deep ANN in kcal/mol for test data divided by metal and oxidation state. The number of test cases is indicated in parentheses

Species	RMS Test error	Min. Abs. Test Error	Max. Abs. Test Error
Cr(II)	3.3 (86)	0.03	11.2
Cr(III)	2.5 (50)	0.06	6.3
Mn(II)	2.8 (57)	0.01	7.5
Mn(III)	2.1 (60)	0.04	5.6
Fe(II)	3.7 (60)	0.07	13.0
Fe(III)	3.0 (71)	0.09	7.9
Co(II)	2.8 (58)	0.07	11.8
Co(III)	4.5 (57)	0.02	15.7
Ni(II)	2.5 (39)	0.09	5.8

Table S25: Average HF exchange sensitivity values, in kcal/mol.HFX, for homoleptic compounds with C, N and O ligands grouped by metal, oxidation state and ligand connecting atom

Meta	I Oxidation state	Ligan	Ligand connectio	
		С	Ν	0
Cr		-82	-55	-59
	111	-20	-20	-18
Mn	II	-167	-92	-87
	111	-58	-44	-32
Fe	II	-164	-76	-59
	111	-118	-74	-63
Co	II	-106	-48	-45
		-98	-53	-46
Ni	II	-65	-57	-52



Figure S9: Scatter plot of HFX sensitivity in kcal/mol \cdot HFX for homoleptic M(II) complexes, colored by connection atom, for homoleptic (II) complexes with C (gray), N (blue) and O (square) ligands.



Figure S10: Error boxplot for regression of $\frac{\partial \Delta E_{H-L}}{\partial a_{HF}}$ using an ANN, showing training and test data comparison. The top number indicates the number of trials, while the bottom indicates the RMSE.


Figure S11: Model predictions of $\frac{\partial \Delta E_{H-L}}{\partial a_{HF}}$ and data for Co using an ANN. The ligands are described by two numbers indicating the equatorial first and then the axial, color coded by ligand identity (green for halogen, gray for carbon, blue for nitrogen, and red for oxygen). The error bars represent an estimated 1 standard deviation from the mean prediction.



Figure S12: Model predictions of $\frac{\partial \Delta E_{H-L}}{\partial a_{HF}}$ and data for Cr using an ANN. The ligands are described by two numbers indicating the equatorial first and then the axial, color coded by ligand identity (green for halogen, gray for carbon, blue for nitrogen, and red for oxygen). The error bars represent an estimated 1 standard deviation from the mean prediction.



Figure S13: Model predictions of $\frac{\partial \Delta E_{H-L}}{\partial a_{HF}}$ and data for low-spin Fe(II) (top) and Ni(II) (bottom) using an ANN. The ligands are described by two numbers indicating the equatorial first and then the axial, color coded by ligand identity (green for halogen, gray for carbon, blue for nitrogen, and red for oxygen). The error bars represent an estimated 1 standard deviation from the mean prediction.



Figure S14: Model predictions of $\frac{\partial \Delta E_{H-L}}{\partial a_{HF}}$ and data for Mn using an ANN. The ligands are described by two numbers indicating the equatorial first and then the axial, color coded by ligand identity (green for halogen, gray for carbon, blue for nitrogen, and red for oxygen). The error bars represent an estimated 1 standard deviation from the mean prediction.

Table S26: HFX sensitivity predictions and data compared in kcal/mol·HFX, compared for the full set of complexes. Each complex is named [metal oxidation axial ligand eq equatorial ligand]. For each complex, the first row has the data obtained from DFT and the second row has the ANN predictions

name	type	HFX sensitivity
cr 2 eq h2o ax ncs	data	-22.2
cr 2 eq porphyrin av nh3	data	-28.1
ci z eq porpriyrin ax mis	pred	-54.4
cr 2 eq scn ax pisc	data	-86.8
	pred	-73
cr 2 eq nos ax pisc	oala	-80.7
cr 2 eg cn ax nh3	data	-54
	pred	-48.4
cr 2 eq tbuc ax tbuc	data	-53.3
cr 2 eg phen ax scn	data	-82.8
	pred	-85.2
cr 2 eq en ax co	data	-81.3
cr 2 eg bipy ax bipy	data	-66.5
-	pred	-60.7
cr 2 eq en ax en	data	-29.2
cr 2 eq co ax cn	data	-83.2
	pred	-76.8
cr 2 eq pisc ax pisc	data	-87.6
cr 2 eg ox ax h2o	data	-59
	pred	-57.7
cr 2 eq h2o ax cl	data	-14.3
cr 2 eq nh3 ax scn	data	-32.5 -31.3
	pred	-58
cr 2 eq ox ax ox	data	-61.6
cr 2 eg h2o ax h2o	data	-53.8 -29.7
	pred	-34.4
cr 2 eq cn ax h2o	data	-18.1
cr 2 eq pisc av pcs	data	-53.8
	pred	-61.2
cr 2 eq acac ax acac	data	-84.7
cr 2 eq bipy ax h2o	data	-53.2 -49
	pred	-62.5
cr 2 eq nh3 ax nh3	data	-28.4
cr 2 eq co ax co	data	-43.8 -84.2
	pred	-87.5
cr 2 eq c2h3ns ax c2h3ns	data	-80.7
cr 2 eq phen ax phen	data	-88.7
or 2 of prior ax prior	pred	-58.1
cr 2 eq tbuc ax nh3	data	-44.9
cr 2 eq nh3 ax cn	data	-50.2 -42.1
	pred	-57.2
cr 3 eq c2h3ns ax c2h3ns	data	-21.9
cr 3 eq acac ax acac	data	-45.5
	pred	-20.7
cr 3 eq h2o ax scn	data	-24.5
cr 3 eg en ax nh3	data	-22.3
	pred	-19.3
cr 3 eq scn ax scn	data	-22.4
cr.3 eg en ax en	data	-27.7
	pred	-19
cr 3 eq cn ax cn	data	-20.1
cr.3 eq.cn.ax.ncs	data	-37.5 -21 4
	pred	-25.1
cr 3 eq tbuc ax tbuc	data	-17.6
or 3 eq nos av nos	pred	-21.6
	pred	-22.7
cr 3 eq ox ax ox	data	-16.6
cr 3 eg cl av cl	pred	-21.7
	pred	-54.4
cr 3 eq co ax co	data	-17.9

	pred	-53.3	
cr 3 eg h2o ax co	data	-22.7	
	pred	-20.8	
cr 3 eq bipy ax bipy	data	-16.1	
	pred	-24.9	
cr 3 eq ncs ax cl	data	-22.2	
or 0 og thug ov pigg	pred	-31.4	
cr 3 eq touc ax pisc	oala	-31.0	
cr 3 eq ox ax cl	data	-16.1	
	pred	-25.9	
cr 3 eg h2o ax h2o	data	-17.8	
	pred	-14.9	
cr 3 eq ox ax pisc	data	-32.3	
	pred	-28.2	
min 2 eq acac ax mis	nred	-110	
mn 2 eg nh3 av nh3	data	-70.3	
	pred	-72.1	
mn 2 eq co ax co	data	-170	
	pred	-140	
mn 2 eq cn ax nh3	data	-94.1	
	pred	-93.2	
min 2 eq en ax en	nred	-01.4	
mn 2 eg acac ax acac	data	-110	
<u>-</u> oq abab ak abab	pred	-103	
mn 2 eq en ax cn	data	-104	
	pred	-109_	
mn 2 eq bipy ax h2o	data	-91.5	_
mn 2 og h 20 og h 20	data	-90.3	
1111 2 EY 1120 at 1120	pred	-63.7	
mn 2 eg ncs ax nh3	data	-81.8	
	pred	-90.3	
mn 2 eq bipy ax bipy	data	-121	
0	pred	-112	
mn 2 eq ox ax ox	oala	-89.1	
mn 2 eg c2h3ns ax c2h3ns	data	-164	
	pred	-141	
mn 2 eq ncs ax co	data	-161	
0	pred	-131	
mn 2 eq ncs ax ncs	data	-94.2	
mn 2 eg phen ax phen	data	-119	
	pred	-110	
mn 2 eq nh3 ax co	data	-147	
	pred	-107	
mn 2 eq en ax n2o	data	-64.6	_
mn 2 eg en ax co	data	-156	
	pred	-118	
mn 2 eq nh3 ax pisc	data	-117	
	pred	-105	
mn 3 eq en ax co	data	-64.8	
mn 3 eg phen av cl	data	-46.1	
min o eq phen ax ci	pred	-57.5	
mn 3 eg phen ax phen	data	-43.5	
	pred	-56.9	
mn 3 eq cn ax nh3	data	-45.7	
mn 3 eg pisc av pisc	data	-90.5	
min 5 eq pisc ax pisc	nred	-95.3	
mn 3 eg ncs ax cl	data	-47.1	
·	pred	-54.3	
mn 3 eq h2o ax co	data	-43.3	
	data	-45.8	
	pred	-43.9	
mn 3 eq co ax co	data	-59.3	
	pred	-89.7	
mn 3 eq acac ax acac	data	-42.9	_
mn 3 eg nh3 ey nh3	data	-48.6	
IIII O EY IIIO AX IIIO	pred	-41.4	
mn 3 eq nh3 ax co	data	-58.5	
	pred	-61.7	
mn 3 eq cn ax cn	data	-50.7	
mn 2 og nh2 ov oon	pred	-80.9	
min o eq milo ax sch	nred	-66.8	
mn 3 eg bipv ax bipv	data	-42.4	
	pred	-58.8	
mn 3 eq h2o ax pisc	data	-137	
mn 3 eg en av en	data	-40.1	
	pred	-42.8	
mn 3 eg phen ax nh3	data	-36.5	
		-	

	pred	-50.2	
1111 3 eq 1120 ax 1120	pred	-28.4	
mn 3 eq c2h3ns ax c2h3ns	data	-62.7	
fe 2 eg co ax co	data	-00.0 -174	
(pred	-144	
te 2 eq acac ax acac	oata	-117	
fe 2 eq tbuc ax co	data	-73	
fe 2 eg cn ax cn	data	-97.6	
	pred	-135	
le 2 eq touc ax touc	pred	-71	
fe 2 eq h2o ax ncs	data	-52.2	
fe 2 eq en ax en	data	-74.5	
fe 2 eg cl av pisc	pred	-86.7	
	pred	-120	
te 2 eq ncs ax h2o	data	-69.4 -81.7	
fe 2 eq pisc ax cl	data	-133	
fe 2 eg phen ax phen	data	-131 -118	
fo 2 og oon ov b2o	pred	-108	
le 2 eq sch ax h20	pred	-66.4	
fe 2 eq nh3 ax nh3	data	-68.5	
fe 2 eq c2h3ns ax c2h3ns	data	-160	
fe 2 eq ox av pisc	pred	-142	
	pred	-107	
te 2 eq pisc ax ch	data pred	-155 -144	
fe 2 eq acac ax cn	data	-104	
fe 2 eq ox ax ox	data	-73.2	
fo 2 og h2o av h2o	pred	-80	
	pred	-68.4	
te 2 eq ncs ax pisc	data	-86.5 -127	
fe 2 eq en ax co	data	-141	
fe 2 eg pisc ax ncs	data	-111 -129	
fo Q og pog ov og	pred	-122	
le 2 eq fics ax co	pred	-124	
fe 2 eq ncs ax ncs	data	-86.4	
fe 3 eq tbuc ax tbuc	data	-61	
fe 3 eg ch ax ch	pred data	-61.7 -112	
	pred	-107	
te 3 eq acac ax acac	data pred	-70.3 -72.4	
fe 3 eq porphyrin ax h2o	data	-65.3	
fe 3 eg h2o ax cn	data	-70.1	
fo 2 og porphyrin av on	pred	-56.7	
ie s eq porpriyrin ax ch	pred	-94.1	
fe 3 eq co ax co	data	-113 -113	
fe 3 eq ox ax ox	data	-65.7	
fe 3 eq pisc ax h2o	data	-63.8 -102	
fo 2 og oppo ov ol	pred	-86	
le 5 eq acac ax ci	pred	-62.9	
fe 3 eq ox ax ncs	data	-72.3	
fe 3 eq en ax en	data	-67.1	
fe 3 eg pisc ax pisc	data	-62.2 -133	
fo 2 og biov ov og	pred	-122	
ie o eq bipy ax co	pred	-122	
fe 3 eq pisc ax ncs	data	-94.5	
fe 3 eq ncs ax cl	data	-81.9	
fe 3 eg phen ax phen	pred data	-79 -87.8	
fo 2 og b2o ov b2c	pred	-82.6	
ie 3 eq nzo ax nzo	pred	-53.2 -42.6	
fe 3 eq cl ax cl	data	-69.6	

		CO 1	
fe 3 eq pisc ax cl	data	-69.1	
	pred	-94.9	
te 3 eq c2h3ns ax c2h3ns	data	-112	
fe 3 eq nh3 ax scn	data	-64.4	
fo 2 og apap av pos	pred	-80.2	
le 5 eq acac ax nos	pred	-61.1	
fe 3 eq co ax scn	data	-90.1	
fe 3 eg bipy ax bipy	data	-102	
	pred	-85.7	
te 3 eq ncs ax cn	data	-75.3 -108	
fe 3 eq ncs ax ncs	data	-62.5	
fe 3 eq.nh3 ay.nh3	data	-75.4	
	pred	-57.8	
fe 3 eq cl ax pisc	data	-77.8	
co 2 eq ox ax ox	data	-41.8	
	pred	-52.4	
	pred	-56.2	
co 2 eq tbuc ax tbuc	data	-36	
co 2 eg cl ax pisc	data	-47.9	
	pred	-76.1	
co 2 eq cn ax n2o	pred	-52.9 -57	
co 2 eq en ax pisc	data	-78.6	
co 2 eg phen ax h2o	data	-82.4 -59.4	
	pred	-63.5	
co 2 eq phen ax phen	data	-65.8 -74.3	
co 2 eq h2o ax ncs	data	-34.6	
co 2 eq c2h3ns ax c2h3ns	pred data	-40.7	
	pred	-102	
co 2 eq nh3 ax nh3	data	-50.4	
co 2 eq acac ax cn	data	-63.1	
co 2 eq nh3 ax h2o	data	-85.3	
	pred	-42.4	
co 2 eq acac ax acac	data	-39.6	
co 2 eq ncs ax ncs	data	-41.9	
co 2 eg bipy ax bipy	pred	-66.7	
	pred	-76.1	
co 2 eq co ax co	data	-109	
co 2 eq en ax h2o	data	-56.3	
co 2 ca h 2 c a x h 2 c	pred	-50.9	
	pred	-35.7	
co 3 eq co ax nh3	data	-98.4	
co 3 eq acac ax nh3	data	-42.8	
an 2 ng thun ay ping	pred	-52.4	
co s eq ibuc ax pisc	pred	-74.4	
co 3 eq nh3 ax nh3	data	-60.4	
co 3 eq ox ax ox	data	-68.6	
	pred	-53.1	
co s eq en ax en	pred	-58.1	
co 3 eq acac ax acac	data	-47.5	
co 3 eq phen ax phen	data	-၁၁.୪ -76	
	pred	-72.5	
co 3 eq ch ax ch	pred	-95.3 -89	
co 3 eq c2h3ns ax c2h3ns	data	-87.7	
co 3 eg co ax co	data	-92.7	
	pred	-94.2	
co 3 eq tbuc ax tbuc	data	-31./ -48.7	
co 3 eq h2o ax h2o	data	-25.3	
co 3 eq nos av nos	pred	-36.8 -23.2	
	pred	-67.6	
co 3 eq pisc ax pisc	data	-/6.4 -94.8	
co 3 eq bipy ax bipy	data	-74	

	pred	-73.2	
co 3 eq acac ax ncs	data	-61.9	
	pred	-54.8	
co 3 eq ox ax ncs	data	-66	
	pred	-50.5	
co 3 eq acac ax co	data	-86	
	pred	-75.8	
ni 2 eq tbuc ax tbuc	data	-60.6	
	pred	-59.8	
ni 2 eq bipy ax nh3	data	-57.9	
	pred	-59.3	
ni 2 eq en ax en	data	-56.1	
	pred	-54.8	
ni 2 eq co ax co	data	-67.2	
	prea	-79.3	
ni 2 eq nn3 ax nn3	data	-73.1	
	prea	-55.2	
ni 2 eq acac ax cn	data	-60	
ni 0 og pige ev pige	pred	-60.4	
ni 2 eq pisc ax pisc	uala	-80.4	
ni 2 og op ov b2o	dete	-00.1	
ni z eq en ax nzo	uala	-01.4	
ni 2 eq hiny ay hiny	data	-62.7	
The 2 eq bipy as bipy	nred	-60.7	
ni 2 eq nos av nos	data	-53.2	
	nred	-59.5	
ni 2 eg ch av ch	data	-65.6	
	pred	-69.8	
ni 2 eq acac ax co	data	-52.4	
	pred	-63.1	
ni 2 eq c2h3ns ax c2h3ns	data	-62.1	
	pred	-76.6	
ni 2 eq acac ax nh3	data	-50.4	
	pred	-51.1	
ni 2 eq acac ax acac	data	-52.4	
	pred	-55	
ni 2 eg ox ax ox	data	-69.2	
	nred	-59 5	



Figure S15: Parity plot for 2 standard deviation from the mean prediction and absolute prediction error for test case $\frac{\partial \Delta E_{H-L}}{\partial a_{HF}}$ prediction using ANN. All units are kcal/mol.HFX. The black line is y=x.



Figure S16: Error boxplot for regression of R_{LS}^{min} using an ANN, showing training and test data comparison. The top number indicates the number of trials, while the bottom indicates the RMSE.



Figure S17: Parity plot for 2 standard deviation from the mean prediction and absolute prediction error for test case R_{LS}^{min} prediction using ANN. All units are Å. The black line is y=x.



Figure S18: Model predictions of R_{LS}^{min} and data for low-spin Co using an ANN. The ligands are described by two numbers indicating the equatorial first and then the axial, color coded by ligand identity (green for halogen, gray for carbon, blue for nitrogen, and red for oxygen). The error bars represent an estimated 1 standard deviation from the mean prediction.



Figure S19: Model predictions of R_{LS}^{min} and data for low-spin Cr using an ANN. The ligands are described by two numbers indicating the equatorial first and then the axial, color coded by ligand identity (green for halogen, gray for carbon, blue for nitrogen, and red for oxygen). The error bars represent an estimated 1 standard deviation from the mean prediction.



Figure S20: Model predictions of R_{LS}^{min} and data for low-spin Fe(II) (top) and Ni(II) (bottom) using an ANN. The ligands are described by two numbers indicating the equatorial first and then the axial, color coded by ligand identity (green for halogen, gray for carbon, blue for nitrogen, and red for oxygen). The error bars represent an estimated 1 standard deviation from the mean prediction.



Figure S21: Model predictions of R_{LS}^{min} and data for low-spin Mn using an ANN. The ligands are described by two numbers indicating the equatorial first and then the axial, color coded by ligand identity (green for halogen, gray for carbon, blue for nitrogen, and red for oxygen). The error bars represent an estimated 1 standard deviation from the mean prediction.

Table S27: LS bond distance predictions and data in Å, compared for the full set of complexes. Each complex is named [metal oxidation axial ligand eq equatorial ligand]. For each complex, the first row has the data obtained from DFT using varying fractions of exact exchange (a_{HF}) and the second row has the ANN predictions

	h un n		0.05		a _{HF}	0.0	0.05	
cr 2 eq h2o ax ncs	data	1.98	0.05	2	2.01	2.01	2.02	2.03
	pred	1.97	1.98	1.98	1.99	1.99	2	2.01
cr 2 eq porphyrin ax nh3	data	2.04	2.04	2.04	2.04	2.03	2.03	2.03
cr 2 eq scn ax pisc	data	1.96	1.97	1.97	1.98	1.99	2	2
or 2 og nos av niso	pred	1.97	1.98	1.99	2	2.01	2.03	2.04
ci z eq nes ax pise	pred	1.99	1.97	1.98	1.99	1.99	2.02	2.03
cr 2 eq cn ax nh3	data	2.09	2.08	2.09	2.1	2.11	2.12	2.13
cr 2 eq tbuc ax tbuc	data	2.06	2.07	2.08	2.08	2.09	2.1	2.11
	pred	1.99	1.99	2	2	2	2	2.01
cr 2 eq phen ax sch	data	2.07	2.07	2.08	2.07	2.07	2.08	2.09
cr 2 eq en ax co	data	2	2	2.01	2.03	2.04	2.05	2.07
cr 2 eq hiny ay hiny	pred	1.98	2 04	2.02	2.03	2.04	2.06	2.07
	pred	2.05	2.05	2.05	2.04	2.06	2.07	2.07
cr 2 eq en ax en	data	2.18	2.19	2.19	2.19	2.19	2.19	2.2
cr 2 eq co ax cn	data	2.17	1.98	2.17	1.99	2.17	2.17	2.18
	pred	2.02	2.03	2.03	2.04	2.04	2.04	2.05
cr 2 eq pisc ax pisc	pred	1.99	2	2.01	2.02	2.03	2.04	2.05
cr 2 eq ox ax h2o	data	1.97	1.96	1.96	1.95	1.95	-	-
or 2 og h2o av ol	pred	1.96	1.96	1.96	1.96	1.97	21	21
	pred	2.09	2.09	2.09	2.09	2.09	2.09	2.09
cr 2 eq nh3 ax scn	data	2.17	2.18	2.18	2.18	2.18	2.18	2.18
cr 2 eq ox ax ox	data	2.17	2.17	2.17	2.10	1.99	1.99	1.98
	pred	1.99	1.99	1.99	2	2	2	2
cr 2 eq n20 ax n20	pred	2.1	2.09	2.09	2.09	2.09	2.09	2.09
cr 2 eq cn ax h2o	data	2.08	2.09	2.1	2.1	2.11	2.11	2.12
cr 2 eq pisc av nos	data	2.06	2.06	2.07	2.07	2.08	2.09	2.09
	pred		1.97	1.97	1.98	1.98	1.99	2
cr 2 eq acac ax acac	data	1.95	1.95	1.95	1.96	1.98	2	2.01
cr 2 eq bipy ax h2o	data	2.08	2.09	2.09	2.09	2.11	2.12	2.12
or 2 og ph2 ov ph2	pred	2.06	2.07	2.08	2.08	2.09	2.1	2.1
	pred	2.2	2.2	2.2	2.21	2.21	2.21	2.22
cr 2 eq co ax co	data	2.05	2.06	2.07	2.08	2.09	2.1	2.11
cr 2 eq c2h3ns ax c2h3ns	data	2.06	2.06	2.07	2.07	2.08	2.08	2.09
	pred	2.02	2.03	2.04	2.04	2.05	2.05	2.06
cr 2 eq pnen ax pnen	pred	2.05	2.05	2.06	2.06	2.07	2.09	2.09
cr 2 eq tbuc ax nh3	data	1.98	1.98	1.98	1.97	1.97	1.97	1.96
cr 2 eq nh3 ax cn	pred data	1.95	1.96	1.96	1.96	1.96	1.97	1.97
	pred	2.04	2.05	2.07	2.08	2.09	2.1	2.11
cr 3 eq c2h3ns ax c2h3ns	data	2.07	2.07	2.08	2.08	2.09	2.09	2.09
cr 3 eq acac ax acac	data	1.96	1.96	1.96	1.96	1.95	1.95	1.96
or 2 og b2o ov son	pred	1.92	1.93	1.93	1.93	1.94	1.94	1.95
ci s eq lizo ax sch	pred	2.05	2.04	2.04	2.03	2.03	2.03	2.02
cr 3 eq en ax nh3	data	2.15	2.14	2.14	2.13	2.13	2.12	2.12
cr 3 eq scn ax scn	data	2.13	2.13	2.13	2.13	2.13	2.13	2.14
	pred	2.46	2.46	2.46	2.46	2.46	2.46	2.46
cr 3 eq en ax en	data	2.14	2.13	2.13	2.13	2.12	2.12	2.12
cr 3 eq cn ax cn	data	2.1	2.11	2.11	2.11	2.11	2.11	2.12
cr 3 eq ch ay nes	pred	2.08	2.08	2.09	2.09	2.1	2.1	2.1
or o eq en ax neo	pred	2.03	2.03	2.04	2.04	2.05		
cr 3 eq tbuc ax tbuc	data	-	-	2.01	2	2	2	2
cr 3 eq ncs ax ncs	data	2.01	2.01	2.01	2.01	2.01	2.02	2.02
	pred	2.01	2.02	2.02	2.02	2.03	2.03	2.03
G S EQ UX AX UX	pred	1.96	1.96	1.96	1.99	1.99	1.99	1.98
cr 3 eq cl ax cl	data	2.5	2.49	2.48	2.49	2.49	2.48	2.48
	prea	∠.40	∠.40	∠.40	∠.40	∠.40	∠.40	∠.40

cr 3 eq co ax co	data	2.15	2.15	2.16	2.16	2.16	2.16	2.17
cr 3 eq h2o ax co	data	1.98	1.98	1.98	1.97	1.97	1.97	1.97
cr 3 eq bipy ax bipy	data	2.07	2.07	2.07	2.07	2.07	2.07	2.07
cr 3 eq ncs ax cl	data	2.06	2.06	2.07	2.07	2.07	2.07	2.07
cr 3 eq tbuc ax pisc	data	2.02	2.02	2.03	2.03	2.04	2.04	2.05
cr 3 eq ox ax cl	pred data	1.93 1.99	1.93 1.99	1.93 1.98	1.94 1.98	1.94 1.97	1.94 1.97	1.94 1.97
cr 3 eq h2o ax h2o	pred data	1.97 2	1.97 2	1.97 1.99	1.97 1.99	1.97 1.99	1.97 1.99	1.98 1.98
cr 3 eq ox ax pisc	pred data	1.98 1.95	1.98 1.95	1.98 1.95	1.98 1.95	1.98 1.95	1.99 1.94	1.99 1.94
mn 2 eg acac ax nh3	pred data	1.93 1.93	1.94 1.94	1.94 1.95	1.94 1.96	1.94 1.98	1.94 2.01	1.94 2.02
mn 2 eq nh3 ax nh3	pred data	1.95 2.15	1.95 2.15	1.95 2.16	1.96 2.16	1.96 2.16	1.97 2.16	1.98 2.16
mn 2 eg co ax co	pred data	2.13 1.98	2.14 1.98	2.15 1.99	2.16 2	2.16 2.01	2.17 2.03	2.17 2.04
mn 2 eg cn ax nh3	pred	1.99	2	2.01	2.01	2.02	2.03	2.04
mn 2 eg en ax en	pred	1.98	2 13	2.01	2.03	2.04	2.05	2.07
	pred	2.11	2.12	2.12	2.13	2.13	2.14	2.14
mn 2 eg en av en	pred	1.93	1.94	1.94	1.95	1.96	1.97	1.98
	pred	1.94	1.95	1.95	1.96	1.98	1.99	2
	pred	2.04	2.05	2.07	2.06	2.08	2.09	2.09
mn 2 eq n2o ax n2o	pred	2.06	2.06	2.06	2.07	2.08	2.09	2.09
mn 2 eq ncs ax nh3	data pred	1.99	2	2.02	2.03	2.04 2.05	2.05	2.06
mn 2 eq bipy ax bipy	data pred	2.02 2.02	2.03 2.03	2.03 2.04	2.05 2.04	2.05 2.05	2.06 2.06	2.06 2.07
mn 2 eq ox ax ox	data pred	2.03 2.03	2.05 2.04	2.08 2.05	2.09 2.05	2.11 2.05	-	-
mn 2 eq c2h3ns ax c2h3ns	data pred	1.96 1.95	1.97 1.96	1.97 1.96	1.98 1.97	1.99 1.98	2 1.98	2.01 1.99
mn 2 eq ncs ax co	data pred	1.86 1.86	1.87 1.86	1.87 1.87	1.88 1.88	1.89 1.89	1.9 1.9	1.91 1.91
mn 2 eq ncs ax ncs	data	2.03	2.04	2.05	2.07	2.08	2.1	2.14
mn 2 eq phen ax phen	data	2.03	2.04	2.05	2.05	2.06	2.05	2.06
mn 2 eq nh3 ax co	data	1.89	1.9	1.91	1.92	1.93	1.95	1.96
mn 2 eq en ax h2o	data	2.09	2.09	2.09	2.1	2.1	2.1	2.1
mn 2 eq en ax co	data	1.88	1.88	1.89	1.9	1.91	1.93	1.94
mn 2 eq nh3 ax pisc	data	1.94	1.95	1.96	1.97	1.99	2.01	2.02
mn 3 eq en ax co	data	2.01	2.02	2.04	2.05	2.06	2.07	2.02
mn 3 eq phen ax cl	data	2.08	2.01	2.02	2.03	2.04	2.04	2.05
mn 3 eq phen ax phen	data	2.07	2.07	2.07	2.08	2.08	2.09	2.02
mn 3 eq cn ax nh3	data	2.03	2.03	2.03	2.04	2.04	2.04	2.04
mn 3 eq pisc ax pisc	data	1.92	2.01	2.01	1.95	2.02	2.02	2.03
mn 3 eq ncs ax cl	pred data	1.92 1.98	1.93	1.94 1.99	1.94 1.99	1.95 1.99	1.95	1.96 2
mn 3 eq h2o ax co	pred data	1.98 1.95	1.99 1.95	1.99 1.94	2 1.94	2.01 1.93	2.02 1.93	2.03 1.93
mn 3 eq ox ax ox	pred data	1.93 1.97	1.94 1.97	1.94 1.96	1.94 1.96	1.94 1.95	1.94 1.95	1.94 1.94
mn 3 eq co ax co	pred data	1.94 2.05	1.94 2.06	1.94 2.06	1.94 2.07	1.94 2.07	1.94 2.08	1.94 2.08
mn 3 eg acac ax acac	pred data	2.06 1.91	2.06 1.92	2.07 1.92	2.08 1.9	2.08 1.92	2.09 1.92	2.09 1.92
mn 3 eq nh3 ax nh3	pred data	1.91 2.11	1.91 2.1	1.91 2.1	1.91 2.09	1.92 2.09	1.92 2.08	1.92 2.08
mn 3 eq nh3 ax co	pred data	2.1	2.1	2.1	2.1	2.1	2.1	2.1
mn 3 eg cn ax cn	pred data	2.02	2.04	2.04	2.05	2.06	2.06	2.07
mn 3 eg nh3 av son	pred	2.03	2.04	2.05	2.05	2.06	2.06	2.07
mn 3 eg hinv av binv	pred	2.09	2.09	2.08	2.08	2.08	2.09	2.09
mn 3 eg blog av blog	pred	2.02	2.02	2.02	2.02	2.02	2.02	2.02
	pred	1.93	1.94	1.94	1.94	1.95	1.95	2.07
nin o eq en ax en	pred	2.08	2.08	2.08	2.08	2.08	2.08	2.07

mn 3 eq phen ax nh3	data	2.06	2.05	2.05	2.04	2.04	2.04	2.04
mn 3 eq h2o ax h2o	data	1.97	1.96	1.96	1.96	1.95	1.95	1.95
mn 3 eq c2h3ns ax c2h3ns	data	1.99	2	2.01	2.01	2.02	2.02	2.02
fe 2 eq co ax co	data	1.92	1.92	1.93	1.93	2.02	1.95	2.04
fe 2 eq acac ax acac	pred data	1.94 1.94	1.95 1.94	1.95 1.95	1.96 1.95	1.97 1.97	1.97 1.97	1.98 1.98
fe 2 eq tbuc ax co	pred data	1.92 1.82	1.93 1.82	1.93 1.83	1.94 1.84	1.94 1.85	1.95 1.86	1.96 1.87
fe 2 eg cn ax cn	pred data	1.85 2	1.85 2	1.85 2.01	1.85	1.85 2.04	1.86	1.86
fe 2 eq thuc ax thuc	pred	1.96	1.97	1.98	1.99	2	2.01	2.02
	pred	2.04	2.04	2.05	2.05	2.05	2.06	1.00
	pred	1.93	1.93	1.94	1.94	1.95	1.95	1.96
te 2 eq en ax en	pred	2.08	2.08	2.09	2.09	2.09	2.09	2.09
te 2 eq cl ax pisc	data pred	1.81 1.85	1.82 1.85	1.82 1.85	1.83 1.85	1.84 1.85	1.86 1.86	1.87 1.86
fe 2 eq ncs ax h2o	data pred	1.93 1.95	1.94 1.95	1.95 1.96	1.98 1.96	1.98 1.97	1.99 1.98	2 1.99
fe 2 eq pisc ax cl	data	1.87	1.87	1.88	1.88	1.89	1.9	1.91
fe 2 eq phen ax phen	data	2.01	2.01	2.02	2.02	2.03	2.03	2.04
fe 2 eq scn ax h2o	data	2.05	2.02	2.03	2.03	2.04	2.03	2.00
fe 2 eq nh3 ax nh3	data	2.04	2.04	2.05	2.05	2.05	2.05	2.05
fe 2 eq c2h3ns ax c2h3ns	pred data	2.05 1.91	2.06 1.91	2.08	2.08	2.09 1.93	2.1 1.94	2.1 1.94
fe 2 eq ox ax pisc	pred data	1.9 1.83	1.91 1.83	1.91 1.84	1.91 1.85	1.92 1.86	1.92 1.87	1.93 1.89
fe 2 eq pisc ax cn	pred data	1.85	1.85	1.85	1.86	1.86	1.86	1.87
fo 2 og ploo ax on	pred	1.87	1.87	1.88	1.88	1.88	1.89	1.89
	pred	1.92	1.92	1.92	1.92	1.92	1.92	1.93
te 2 eq ox ax ox	pred	2.03	2.04	2.04	2.04	2.05	2.05	2.05
te 2 eq h2o ax h2o	data pred	2.03 1.99	2.03 2	2.02 2	2.02 2.01	2.02 2.01	2.02 2.01	2.02 2.02
fe 2 eq ncs ax pisc	data pred	1.85 1.85	1.86 1.85	1.87 1.85	1.88 1.86	1.89 1.86	-	1.92 1.88
fe 2 eq en ax co	data	1.85	1.86	1.87	1.88	1.89	1.91 1.9	1.92
fe 2 eq pisc ax ncs	data	1.88	1.89	1.89	1.9	1.91	1.92	1.93
fe 2 eq ncs ax co	data	1.84	1.84	1.85	1.86	1.87	1.88	1.9
fe 2 eq ncs ax ncs	data	1.99	2	2.01	2.03	2.03	2.04	2.05
fe 3 eq tbuc ax tbuc	data	1.98	1.96	1.96	1.95	1.95	2.04 1.95	1.94
fe 3 eq cn ax cn	pred data	1.93 1.98	1.94 1.99	1.94 1.99	1.94 1.99	1.94 2	1.94 2	1.94 2
fe 3 eg acac ax acac	pred data	1.98 1.92	1.99 1.92	1.99 1.92	2 1.91	2 1.91	2.01 1.91	2.01 1.9
fe 3 eq porphyrin ax h2o	pred data	1.91	1.91	1.91	1.91	1.91	1.91	1.91
fe 3 eq h2o ay on	pred	1.99	1.99	1.99	1.99	1.99	2	2
	pred	1.92	1.92	1.92	1.92	1.93	1.93	1.93
te 3 eq porpriyrin ax ch	pred	1.98	1.98	1.98	1.98	1.99	1.99	1.99
te 3 eq co ax co	pred	2.01	2.01	2.02	2.02	2.03	2.03	2.03
te 3 eq ox ax ox	data pred	1.95 1.93	1.95 1.93	1.95 1.93	1.94 1.93	1.94 1.93	1.93 1.93	1.93 1.94
fe 3 eq pisc ax h2o	data pred	1.92 1.91	1.92 1.91	1.93 1.91	1.94 1.91	1.95 1.91	1.96 1.91	1.97 1.91
fe 3 eq acac ax cl	data	1.95	1.95	1.95	1.94	1.94	1.94	1.93
fe 3 eq ox ax ncs	data	1.95	1.95	1.94	1.94	1.93	1.92	1.92
fe 3 eq en ax en	data	2.07	2.07	2.06	2.05	2.05	2.05	2.05
fe 3 eq pisc ax pisc	data	1.91	1.91	1.91	1.92	1.92	1.92	1.94
fe 3 eq bipy ax co	data	1.9 1.89	1.9 1.9	1.9	1.91 1.9	1.91 1.92	1.91 1.93	1.91 1.95
fe 3 eq pisc ax ncs	pred data	1.89 1.9	1.9 1.9	1.91 1.9	1.92 1.89	1.92 1.89	1.93 1.89	1.94 1.88
fe 3 eg ncs ax cl	pred data	1.91 1.94	1.91 1.95	1.91 1.95	1.91 1.95	1.92 1.95	1.92 1.95	1.92 1.96
fe 3 eg phen av phen	pred	1.94	1.94	1.95	1.95	1.96	1.96	1.97
fo 3 og h2c av h2c	pred	2.01	2.01	2.01	2.01	2.01	2.01	2.01
10 0 04 1120 ax 1120	pred	1.93	1.93	1.93	1.93	1.93	1.93	1.93

fe 3 eq cl ax cl	data	2.46	2.46	2.46	2.45	2.45	2.45	2.46
fe 3 eq pisc ax cl	data	1.92	1.93	1.93	1.94	1.94	1.95	1.96
fe 3 eq c2h3ns ax c2h3ns	pred data	1.9 1.95	1.91	1.91	1.92	1.92 1.97	1.93 1.97	1.94 1.98
fe 3 eq nh3 ax scn	data	1.94 2.06	1.95 2.06	1.95 2.05	1.95 2.05	1.96 2.05	1.96	1.96
fe 3 eq acac ax ncs	pred data	2.05 1.88	2.05 1.88	2.05 1.88	2.05 1.88	2.05 1.88	1.88	1.89
fe 3 eq co ax scn	pred data	1.9 -	1.9 -	1.9 1.88	1.9 1.89	1.9 1.9	1.9 1.9	1.9 1.91
fe 3 eg bipy ax bipy	pred data	2	2	1.93	1.93	1.93	1.94	1.94
fe 3 eq nos ax on	pred	2 1 94	2 1.95	2 1 95	2 1.95	2 1.96	2 1.96	2 1.96
fe 3 eq nes av nes	pred	1.91	1.92	1.92	1.93	1.93	1.93	1.94
fo 2 og ph2 ov ph2	pred	1.94	1.94	1.94	1.94	1.95	1.95	1.95
	pred	2.08	2.07	2.07	2.06	2.06	2.06	2.05
ie 3 eq ci ax pisc	pred	1.86	1.87	1.88	1.89	1.9	1.92	1.93
co 2 eq ox ax ox	data pred	2.02 2.01	2.04 2.01	2.04 2.02	2.05 2.03	2.05 2.04	2.06 2.04	2.05 2.05
co 2 eq en ax en	data pred	2.04 2.04	2.04 2.05	2.04 2.06	2.04 2.07	2.04 2.07	2.05 2.08	2.06 2.08
co 2 eq tbuc ax tbuc	data pred	2.06 2.03	2.07 2.04	2.07 2.04	2.08 2.05	2.08 2.05	2.08 2.05	2.05 2.06
co 2 eq cl ax pisc	data	1.82 1.85	1.82 1.85	1.83 1.85	1.84 1.85	1.86 1.85	1.87 1.85	1.89 1.85
co 2 eq cn ax h2o	data	1.94	1.95	1.95	1.96	1.97	1.98	1.98
co 2 eq en ax pisc	data	1.89	1.9	1.91	1.93	1.94	1.96	1.97
co 2 eq phen ax h2o	data	2.02	2.02	2.03	2.03	2.04	2.05	2.06
co 2 eq phen ax phen	data	1.98	1.99	1.99	2.17	2.18	2.18	2.18
co 2 eq h2o ax ncs	data	1.99	1.92	1.91	1.92	1.92	1.93	2.06
co 2 eq c2h3ns ax c2h3ns	data	1.91	1.92	1.92	1.92	1.93	1.93	1.93
co 2 eq nh3 ax nh3	data	1.89 2.04	1.89 2.04	1.9 2.04	1.9 2.04	1.9 2.05	1.91 2.06	1.91 2.07
co 2 eq acac ax cn	pred data	2.01 1.93	2.02 1.93	2.03 1.94	2.05 1.94	2.06 1.95	2.07 1.95	2.07 1.96
co 2 eq nh3 ax h2o	pred data	1.91 2.02	1.91 2.02	1.92 2.02	1.92 2.02	1.92 2.02	1.92 2.02	1.93 2.02
co 2 eq acac ax acac	pred data	2 1.92	2 1.93	2.01 1.94	2.01 1.96	2.02 1.96	2.02 1.97	2.03 1.97
co 2 eq ncs ax ncs	pred data	1.93 1.93	1.93 1.96	1.93 1.97	1.94 1.99	1.94 2	1.95 2.01	1.95 2.02
co 2 eq bipy ax bipy	pred data	1.92	1.94	1.96	1.97	1.99 1.99	2	2.02
$c_0 2 eq c_0 ax c_0$	pred	1.99	2	2.01	2.02	2.03	2.04	2.05
co 2 eq en ax b2o	pred	1.91	1.92	1.92	1.93	1.93	1.94	1.95
	pred	2.01	2.02	2.02	2.03	2.03	2.04	2.04
	pred	1.99	1.99	2.01	2.01	2.01	2.01	2.01
co 3 eq co ax nn3	pred	-	1.94	1.95	1.96	1.96	1.97	1.98
co 3 eq acac ax nh3	data pred	1.93 1.9 <u>1</u>	1.93	1.92	1.92	1.91	1.91	1.9 1.91
co 3 eq tbuc ax pisc	data pred	1.87 1.87	1.88 1.88	1.89 1.88	1.89 1.88	1.9 1.88	1.91 1.89	1.91 1.89
co 3 eq nh3 ax nh3	data pred	2.05 2.03	2.04 2.03	2.04 2.03	2.03 2.03	2.03 2.04	2.02 2.04	2.02 2.04
co 3 eq ox ax ox	data pred	1.87 1.92	1.88 1.92	1.88 1.92	1.87 1.93	1.87 1.93	1.87 1.93	1.86 1.93
co 3 eq en ax en	data	2.04	2.03	2.03	2.02	2.02	2.02	2.01
co 3 eq acac ax acac	data	1.93	1.93	1.92	1.92	1.91	1.9	1.9
co 3 eq phen ax phen	data	1.99	1.99	1.99	1.98	1.98	1.98	1.98
co 3 eq cn ax cn	data	1.95	1.95	1.95	1.95	1.95	1.96	1.96
co 3 eq c2h3ns ax c2h3ns	data	1.94	1.95	1.95	1.90	1.90	1.97	1.90
co 3 eq co ax co	data	1.92	1.92	1.93	1.93	1.93	1.94	1.94
co 3 eq tbuc ax tbuc	pred data	1.96	1.96	1.97	1.97 1.94	1.98	1.98	1.99
co 3 eq h2o ax h2o	pred data	1.93	1.93 1.95	1.93 1.95	1.93	1.94 1.93	1.94 1.92	1.94 1.92
co 3 eq ncs ax ncs	pred data	1.93 1.92	1.93 1.92	1.93 1.92	1.93 1.92	1.93 1.92	1.93 1.92	1.93 1.92
co 3 eq pisc ax pisc	pred data	1.92 1.91	1.92 1.91	1.92 1.91	1.92 1.91	1.93 1.91	1.93 1.91	1.93 1.92
	pred	1.88	1.89	1.89	1.89	1.9	1.9	1.9

co 3 eq bipy ax bipy	data	1.98	1.98	1.98	1.98	1.97	1.97	1.97
1 1 1	pred	1.99	1.99	1.99	1.99	2	2	2
co 3 eg acac ax ncs	data	1.93	1.93	1.92	1.92	1.92	1.91	1.91
	pred	1.9	1.9	1.9	1.9	1.9	1.9	1.9
co 3 eq ox ax ncs	data	1.94	1.94	1.94	1.93	1.92	1.91	1.91
•	pred	1.9	1.9	1.9	1.9	1.91	1.91	1.91
co 3 eq acac ax co	data	1.93	1.94	1.94	1.93	1.92	1.91	1.91
	pred	1.91	1.91	1.91	1.91	1.91	1.91	1.91
ni 2 eq tbuc ax tbuc	data	1.93	1.93	1.92	1.92	1.91	1.91	1.91
•	pred	1.91	1.91	1.91	1.92	1.92	1.92	1.93
ni 2 eq bipy ax nh3	data	1.95	1.95	1.95	1.95	1.95	1.96	1.95
	pred	1.94	1.94	1.94	1.94	1.94	1.95	1.96
ni 2 eq en ax en	data	1.98	1.98	1.98	1.98	1.98	1.98	1.98
•	pred	1.96	1.97	1.97	1.97	1.98	1.98	1.99
ni 2 eq co ax co	data	1.87	1.88	1.89	1.9	1.91	1.92	1.93
•	pred	1.9	1.9	1.9	1.91	1.91	1.92	1.92
ni 2 eg nh3 ax nh3	data	1.95	1.95	1.95	1.95	1.95	1.95	1.95
•	pred	1.93	1.94	1.94	1.94	1.95	1.96	1.96
ni 2 eq acac ax cn	data	1.9	1.9	1.9	1.91	1.91	1.91	1.92
•	pred	1.89	1.89	1.89	1.89	1.89	1.9	1.9
ni 2 eq pisc ax pisc	data	1.87	1.87	1.88	1.88	1.89	1.89	1.9
	pred	1.86	1.86	1.86	1.87	1.87	1.87	1.87
ni 2 eq en ax h2o	data	1.97	1.97	1.97	1.97	1.96	1.96	1.96
•	pred	1.96	1.96	1.96	1.96	1.96	1.96	1.96
ni 2 eq bipy ax bipy	data	1.94	1.95	1.95	1.95	1.95	1.96	1.96
	pred	1.94	1.94	1.95	1.95	1.96	1.96	1.97
ni 2 eq ncs ax ncs	data	1.88	1.88	1.88	1.89	1.89	1.9	1.9
	pred	1.87	1.87	1.87	1.88	1.89	1.89	1.9
ni 2 eq cn ax cn	data	-	-	1.94	1.94	1.95	1.96	1.96
	pred			1.92	1.92	1.93	1.93	1.94
ni 2 eq acac ax co	data	1.89	1.89	1.88	1.88	1.88	1.88	1.88
	pred	1.87	1.87	1.87	1.88	1.88	1.88	1.88
ni 2 eq c2h3ns ax c2h3ns	data	1.86	1.87	1.87	1.88	1.89	1.89	1.9
•	pred	1.88	1.88	1.88	1.89	1.89	1.89	1.89
ni 2 eq acac ax nh3	data	1.95	1.94	1.94	1.94	1.94	1.94	1.94
	pred	1.91	1.91	1.91	1.91	1.91	1.91	1.91
ni 2 eq acac ax acac	data	1.89	1.89	1.89	1.89	1.89	1.89	1.9
	pred	1.9	1.9	1.9	1.9	1.9	1.9	1.9
ni 2 eq ox ax ox	data	1.93	1.92	1.92	1.91	1.91	1.91	1.91
•	pred	1.91	1.91	1.91	1.91	1.91	1.91	1.92



Figure S22: Error boxplot for regression of R_{HS}^{min} using an ANN, showing training and test data comparison. The top number indicates the number of trials, while the bottom indicates the RMSE.



Figure S23: Parity plot for 2 standard deviation from the mean prediction and absolute prediction error for test case R_{HS}^{min} prediction using ANN. All units are Å. The black line is y=x.



Figure S24: Model predictions of R_{HS}^{min} and data for high-spin Co using an ANN. The ligands are described by two numbers indicating the equatorial first and then the axial, color coded by ligand identity (green for halogen, gray for carbon, blue for nitrogen, and red for oxygen). The error bars represent an estimated 1 standard deviation from the mean prediction.



Figure S25: Model predictions of R_{HS}^{min} and data for high-spin Cr using an ANN. The ligands are described by two numbers indicating the equatorial first and then the axial, color coded by ligand identity (green for halogen, gray for carbon, blue for nitrogen, and red for oxygen). The error bars represent an estimated 1 standard deviation from the mean prediction.



Figure S26: Model predictions of R_{HS}^{min} and data for high-spin Fe(II) (top), Fe(III) (middle) and Ni(II) (bottom) using an ANN. The ligands are described by two numbers indicating the equatorial first and then the axial, color coded by ligand identity (green for halogen, gray for carbon, blue for nitrogen, and red for oxygen). The error bars represent an estimated 1 standard deviation from the mean prediction.



Figure S27: Model predictions of R_{HS}^{min} and data for high-spin Mn using an ANN. The ligands are described by two numbers indicating the equatorial first and then the axial, color coded by ligand identity (green for halogen, gray for carbon, blue for nitrogen, and red for oxygen). The error bars represent an estimated 1 standard deviation from the mean prediction.

Table S28: HS bond distance predictions and data in Å, compared for the full set of complexes. Each complex is named [metal oxidation axial ligand eq equatorial ligand]. For each complex, the first row has the data obtained from DFT using varying fractions of exact exchange (a_{HF}) and the second row has the ANN predictions

name					a _{HF}			
name	type	0	0.05	0.1	0.15	0.2	0.25	0.3
cr 2 eq h2o ax ncs	data	1.98	1.98	1.99	2	2	2	2.01
cr 2 eq porphyrin ax nh3	data	2.07	2.07	2.07	2.07	2.05	2.05	2.03
	pred	2.07	2.06	2.06	2.06	2.06	2.06	2.06
cr 2 eq scn ax pisc	data	2.04	2.05	2.07	2.08	2.09	2.1	2.11
	pred	2.12	2.13	2.14	2.15	2.17	2.19	2.21
cr 2 eq rics ax pisc	nred	2.02	2.04	2.04	2.05	2.06	2.06	2.07
cr 2 eg cn ax nh3	data	2.12	2.13	2.14	2.15	2.16	2.19	2.2
•·· _ ••] •·· •·· ···•	pred	2.12	2.12	2.12	2.12	2.13	2.13	2.13
cr 2 eq tbuc ax tbuc	data	2.04	2.05	2.05	2.05	2.06	2.06	2.06
cr 2 eq phen av son	data	2.07	2.07	2.07	2.07	2.07	2.07	2.07
ci z eq pileri ax scri	pred	2.13	2.14	2.14	2.14	2.15	2.16	2.16
cr 2 eq en ax co	data	2.16	2.16	2.16	2.16	2.16	2.16	2.16
	pred	2.16	2.16	2.16	2.16	2.16	2.16	2.16
cr 2 eq bipy ax bipy	oata	2.1	2.12	2.12	2.13	2.13	2.13	2.14
cr 2 eg en ax en	data	2.2	2.2	2.2	2.2	2.2	2.2	2.2
	pred	2.19	2.19	2.19	2.19	2.18	2.18	2.18
cr 2 eq co ax cn	data	2.02	2.04	2.05	2.05	2.06	2.06	2.07
or 2 og nice av nice	pred	2.07	2.07	2.08	2.08	2.09	2.11	2.12
01 2 eg pist at pist	pred	2.11	2.12	2.13	2.14	2.13	2.10	2.15
cr 2 eq ox ax h2o	data	2.01	2.01	2.01	2.01	2.01	2.01	2.01
	pred	2	2	2.01	2.01	2.01	2.02	2.02
cr 2 eq h2o ax cl	data	2.05	2.05	2.05	2.05	2.05	2.05	2.05
cr 2 eq nh3 ax scn	data	2.00	2.00	2.00	2.00	2.00	2.00	2.00
	pred	2.17	2.17	2.17	2.18	2.18	2.18	2.18
cr 2 eq ox ax ox	data	2.07	2.07	2.07	2.08	2.08	2.08	2.08
	pred	2.07	2.07	2.07	2.07	2.07	2.08	2.08
cr 2 eq 1120 ax 1120	nred	2.09	2.09	2.09	2.09	2.09	2.09	2.09
cr 2 eg cn ax h2o	data	2.14	2.15	2.16	2.17	2.17	2.18	2.19
	pred	2.12	2.13	2.13	2.13	2.14	2.14	2.14
cr 2 eq pisc ax ncs	data	-	1.99	2	2	2.01	2.01	2.01
cr 2 eq acac ax acac	data	2 04	2.01	2.01	2.02	2.02	2.02	2.02
	pred	2.05	2.05	2.05	2.05	2.05	2.05	2.05
cr 2 eq bipy ax h2o	data	2.12	2.13	2.13	2.13	2.14	2.14	2.14
	pred	2.14	2.14	2.13	2.13	2.13	2.13	2.13
cr 2 eq nn3 ax nn3	oala	2.21	2.21	2.21	2.21	2.22	2.21	2.21
cr 2 eq co ax co	data	2.15	2.16	2.19	2.19	2.2	2.21	2.22
	pred	2.16	2.17	2.18	2.18	2.19	2.2	2.2
cr 2 eq c2h3ns ax c2h3ns	data	2.1	2.11	2.13	2.14	2.14	2.15	2.16
cr 2 eq phen ax phen	data	2.11	2.11	2.12	2.12	2.13	2.14	2.14
or z eq prior ax prior	pred	2.14	2.14	2.14	2.14	2.14	2.14	2.15
cr 2 eq tbuc ax nh3	data	1.99	1.99	2	1.99	2	1.99	2
or 2 og ph2 ov op	pred	1.99	1.99	1.99	1.99	1.99	1.99	1.99
or 2 eq milo ax cm	pred	2.09	2.12	2.11	2.12	2.12	2.13	2.13
cr 3 eq c2h3ns ax c2h3ns	data	2.09	2.09	2.09	2.1	2.1	2.1	2.11
	pred	2.09	2.09	2.1	2.1	2.1	2.11	2.11
cr 3 eq acac ax acac	oata	1.97	1.97	1.97	1.96	1.96	1.96	1.97
cr 3 eq h2o ax scn	data	2.04	2.03	2.03	2.02	2.02	2.02	2.01
	pred	2	2	2	2	1.99	1.99	1.99
cr 3 eq en ax nh3	data	2.15	2.15	2.14	2.14	2.13	2.13	2.13
cr 3 eq scn av scn	data	2.15	2.15	2.15	2.15	2.14	2.14	2.14
	pred	2.51	2.51	2.51	2.51	2.51	2.51	2.51
cr 3 eq en ax en	data	2.16	2.15	2.15	2.14	2.14	2.13	2.13
0	pred	2.15	2.15	2.14	2.14	2.13	2.13	2.13
cr 3 eq cri ax ch	oala	2.12	2.12	2.12	2.13	2.13	2.13	2.13
cr 3 eq cn ax ncs	data	2.05	2.05	2.06	2.06	2.06	2.06	2.07
	pred	2.04	2.05	2.05	2.06	2.07	2.07	2.08
cr 3 eq tbuc ax tbuc	data	-	-	2.02	2.01	2.01	2	2
cr 3 eq nes av nes	dete	2.02	2.02	2.02	2.02	2.02	2.01	2.01
or o eq nos ax nos	pred	2.02	2.02	2.02	2.02	2.02	2.02	2.04
cr 3 eq ox ax ox	data	2.01	2.01	2	2	1.99	1.99	1.99
	pred	2	2	1.99	1.99	1.99	1.99	1.99
cr 3 eq cr ax cr	pred	2.52	2.52	2.51	2.51	2.0	2.0	2.49
	P100							

cr 3 eq co ax co	data	2.17	2.17	2.17	2.18	2.18	2.18	2.18
cr 3 eq h2o ax co	data	2.10	2.10	2.10	1.99	1.99	1.99	1.99
cr 3 eq bipy ax bipy	data	2.08	2.08	2.08	2.08	2.07	2.07	2.07
cr 3 eq ncs ax cl	pred data	2.09 2.03	2.09 2.03	2.09 2.04	2.09 2.04	2.09 2.04	2.08 2.04	2.08 2.04
cr 3 eq tbuc ax pisc	pred data	2.06 1.98	2.06 1.98	2.07 1.97	2.07 1.97	2.08 1.96	2.09 1.95	2.1 1.95
cr 3 eq ox ax cl	pred	1.97	1.96	1.96	1.96	1.96	1.96	1.96
	pred	1.99	1.99	2	2	2	2	2
	pred	2	2	2	2	1.99	1.99	1.99
cr 3 eq ox ax pisc	pred	1.97 1.95	1.96	1.96	1.95	1.95	1.95	1.94 1.94
mn 2 eq acac ax nh3	data pred	2.17 2.16	2.17 2.16	2.17 2.17	2.17 2.17	2.18 2.17	2.18 2.17	2.18 2.17
mn 2 eq nh3 ax nh3	data	2.34	2.34	2.35	2.34 2.31	2.34 2.31	2.34 2.31	2.34 2.31
mn 2 eq co ax co	data	2.41	2.42	2.43	2.43	2.43	2.43	2.43
mn 2 eq cn ax nh3	data	2.17	2.27	2.28	2.29	2.29	2.3	2.31
mn 2 eq en ax en	data	2.20	2.26	2.35	2.28	2.28	2.29	2.3
mn 2 eg acac ax acac	pred data	2.35 2.18	2.35 2.18	2.35 2.18	2.35 2.18	2.34 2.17	2.34 2.17	2.33 2.17
mn 2 eg en ax cn	pred data	2.16	2.16	2.16	2.15	2.15	2.15	2.15
mn 2 eg binv av b2o	pred	2.19	2.19	2.2	2.2	2.2	2.2	2.2
	pred	2.23	2.24	2.24	2.25	2.25	2.25	2.25
	pred	2.16	2.18	2.18	2.18	2.16	2.18	2.16
mn 2 eq ncs ax nh3	data pred	2.16 2.19	2.17 2.19	2.18 2.19	2.17 2.19	2.17 2.19	2.18 2.2	2.18 2.2
mn 2 eq bipy ax bipy	data	2.28	2.28	2.28	2.28	2.28	2.28	2.28
mn 2 eq ox ax ox	data	2.23	2.23	2.23	2.23	2.23	-	-
mn 2 eq c2h3ns ax c2h3ns	data	2.33	2.34	2.34	2.35	2.35	2.35	2.36
mn 2 eq ncs ax co	data	2.15	2.16	2.16	2.16	2.16	2.16	2.16
mn 2 eq ncs ax ncs	data	2.17	2.18	2.18	2.18	2.18	2.18	2.19
mn 2 eq phen ax phen	pred data	2.31 2.3	2.32	2.32	2.32 2.29	2.32 2.29	2.32 2.29	2.32 2.29
mn 2 eg nh3 ax co	pred data	2.29 2.3	2.29 2.3	2.29 2.3	2.28 2.3	2.28 2.29	2.28 2.29	2.28 2.29
mn 2 eg en av h2g	pred	2.28	2.28	2.28	2.28	2.28	2.28	2.28
	pred	2.26	2.26	2.26	2.26	2.26	2.26	2.26
	pred	2.26	2.26	2.26	2.26	2.26	2.26	2.26
min 2 eq mina ax pisc	pred	2.29	2.3	2.31	2.32	2.33	2.34	2.34 2.3 <u>1</u>
mn 3 eq en ax co	data pred	2.09 2.09	2.09 2.09	2.08 2.09	2.08 2.09	2.08 2.09	2.08 2.09	2.07 2.09
mn 3 eq phen ax cl	data pred	2.12 2.17	2.12 2.18	2.24 2.2	2.24 2.21	2.24 2.22	2.24 2.24	2.24 2.25
mn 3 eq phen ax phen	data	2.06	2.06	2.05	2.05	2.05	2.05	2.04
mn 3 eq cn ax nh3	data	2.02	2.04	2.04	2.04	2.04	2.04	2.04
mn 3 eq pisc ax pisc	data	2.03	2.02	2.02	2.03	2.00	2.00	2.00
mn 3 eq ncs ax cl	data	2.02	2.03	2.03	2.03	2.03	2.04	2.04
mn 3 eq h2o ax co	pred data	2.17	2.18 1.92	2.18 1.92	2.19 1.92	2.2 1.91	2.2 1.91	2.21 1.91
mn 3 eg ox ax ox	pred data	1.96 1.95	1.96 1.95	1.95 1.94	1.95 1.94	1.95 1.94	1.95 1.93	1.95 1.93
mn 3 eg co ax co	pred data	1.99 2.12	1.98 2.12	1.98 2.12	1.98 2.12	1.98 2.12	1.98 2.12	1.98 2.12
	pred	2.13	2.13	2.12	2.12	2.13	2.13	2.13
	pred	1.97	1.96	1.96	1.96	1.95	1.95	1.95
mn 3 eq nn3 ax nn3	pred	2.12	2.12	2.12	2.11	2.11	2.1	2.1
mn 3 eq nh3 ax co	data pred	2.11 2.12	2.11 2.11	2.1 2.11	2.1 2.11	2.09 2.1	2.09 2.1	2.09 2.1
mn 3 eq cn ax cn	data pred	2.07 2.11	2.08 2.11	2.08 2.11	2.08 2.11	2.08 2.12	2.05 2.12	2.05 2.13
mn 3 eq nh3 ax scn	data	1.89	1.89	1.91	1.92	1.92 2	1.93	1.94
mn 3 eq bipy ax bipy	data	2.04	2.04	2.04	2.04	2.04	2.04	2.04
mn 3 eq h2o ax pisc	data	2.23	2.23	2.22	2.22	2.21	2.21	-
mn 3 eq en ax en	data	2.11	2.1	2.1	2.09	2.08	2.07	2.08
	pred	2.12	2.11	2.11	2.11	2.1	2.1	2.1

mn 3 eq phen ax nh3	data	2.04	2.03	2.04	2.03	2.03	2.03	2.03
mn 3 eg h2o ax h2o	data	2.07	2.07	2.07	2.07	2.07	2.07	1.95
	pred	1.98	1.98	1.98	1.97	1.97	1.97	1.97
mn 3 eq c2h3ns ax c2h3ns	data pred	2.03	2.03	2.03	2.04	2.04	2.04	2.04
fe 2 eq co ax co	data	2.27	2.29	2.3	2.31	2.31	2.33	2.33
fe 2 eq acac ax acac	data	2.20	2.28	2.29	2.08	2.09	2.09	2.31
fe 2 eq tbuc ax co	data	2.07	2.07	2.08	2.08	2.08	2.08	2.09 2.06
fe 2 eg ch ax ch	pred	1.98 2.44	1.99	2 2 45	2.01	2.02	2.04	2.06
fo 2 og thug ov thug	pred	2.36	2.37	2.38	2.39	2.39	2.39	2.4
	pred	2.2	2.2	2.2	2.2	2.2	2.19 2.19	-
fe 2 eq h2o ax ncs	data pred	1.96 1.99	1.97 1.99	1.98 1.99	1.99 1.99	2 1.99	2.01 2	2.01 2
fe 2 eq en ax en	data	2.27	2.27	2.27	2.27	2.27	2.27	2.27
fe 2 eq cl ax pisc	data	2.01	2.02	2.04	2.05	2.08	2.13	2.15
fe 2 eq ncs ax h2o	data	2.03	2.04 2.05	2.04 2.05	2.05	2.06	2.07	2.09
fe 2 eq pisc ax cl	pred data	2.09	2.09	2.09	2.1 2.21	2.11	2.11	2.12
fo 2 og ploo ux or	pred	2.19	2.2	2.21	2.21	2.22	2.23	2.24
le 2 eq prien ax prien	pred	2.2	2.2	2.2	2.2	2.22	2.02	2.03
fe 2 eq scn ax h2o	data	2.33	2.31	2.3	2.28	2.27	2.29	2.28
fe 2 eq nh3 ax nh3	data	2.31	2.31	2.31	2.31	2.31	2.31	2.32
fo 2 og o2b2no ox o2b2no	pred	2.25	2.25	2.25	2.24	2.24	2.24	2.24
	pred	2.22	2.22	2.23	2.25	2.25	2.20	2.20
te 2 eq ox ax pisc	data pred	2.08 2.02	2.07 2.02	2.07 2.03	2.07 2.03	2.07 2.04	2.08 2.05	2.08 2.06
fe 2 eq pisc ax cn	data	2.14	2.14	2.14	2.14	2.12	2.14	2.14
fe 2 eq acac ax cn	data	2.06	2.07	2.08	2.09	2.08	2.1	2.1
fe 2 eq ox ax ox	data	2.06	2.06	2.07	2.07	2.08	2.08	2.08
fe 2 eq h2o ax h2o	pred data	2.15 2.11						
fe 2 eq nos ax pisc	pred data	2.08	2.08 2.1	2.08 2.1	2.08 2.14	2.09	2.09	2.09
	pred	2.11	2.11	2.11	2.11	2.12		2.12
te 2 eq en ax co	data pred	2.24 2.2	2.23 2.2	2.22 2.2	2.22 2.2	2.22 2.2	2.21 2.2	2.21 2.2
fe 2 eq pisc ax ncs	data	2.05	2.04	2.04	2.04	2.03	2.03	2.03
fe 2 eq ncs ax co	data	2.13	2.12	2.11	2.1	2.09	2.09	2.09
fe 2 eq ncs ax ncs	data	2.23	2.24	2.24	2.24	2.24	2.24	2.25
fe 3 eq tbuc ax tbuc	pred data	2.2 2.09	2.21 2.08	2.22 2.07	2.23 2.06	2.24 2.06	2.25 2.05	2.26 2.05
fe 3 eq ch ax ch	pred data	2.09	2.09	2.08	2.08	2.08	2.07	2.07
	pred	2.24	2.24	2.24	2.23	2.23	2.22	2.22
te 3 eq acac ax acac	pred	2.06	2.05	2.05	2.04	2.03	2.02	2.02
fe 3 eq porphyrin ax h2o	data	2.05	2.05	2.05	2.05	2.05	2.04	2.03
fe 3 eq h2o ax cn	data	2.00	2.00	2.00	2.03	2.03	2.03	2.03
fe 3 eq porphyrin ax cn	data	2.02	2.02	2.02	2.02 2.09	2.02	2.02	2.02
fe 3 eq co ax co	pred data	2.1 2.32	2.1 2.31	2.1 2.31	2.1 2.3	2.1 2.3	2.1 2.29	2.1 2.29
fe 3 eq ox ax ox	pred	2.28	2.28	2.28	2.28	2.28	2.28	2.27
	pred	2.04	2.04	2.03	2.03	2.02	2.02	2.01
te 3 eq pisc ax h2o	data pred	2.16 2.15	2.16 2.15	2.19 2.15	2.18 2.16	2.21 2.15	2.21 2.15	2.2 2.15
fe 3 eq acac ax cl	data	2.14	2.13	2.12	2.11	2.11	2.11	2.1
fe 3 eq ox ax ncs	data	2.03	2.02	2.01	2.01	2	1.99	1.99
fe 3 eq en ax en	data	2.02	2.02 2.25	2.01	2.01	2.01	2.22	2.21
fe 3 eq pisc ax pisc	pred data	2.22 2.17	2.22 2.19	2.21 2.19	2.21 2.2	2.2 2.21	2.2 2.25	2.19 2.23
fe 3 eg hinv av co	pred	2.18	2.19	2.2	2.2	2.21	2.21	2.21
	pred	2.13	2.18	2.17	2.18	2.18	2.18	2.18
te 3 eq pisc ax ncs	data pred	2.05	1.97 2.05	1.96 2.04	1.95 2.04	2.1 2.04	2.1 2.04	2.11 2.05
fe 3 eq ncs ax cl	data pred	2.12 2.14	2.11 2.14	2.11 2.14	2.1 2.14	2.1 2.14	2.1 2.14	2.09 2.14
fe 3 eq phen ax phen	data	2.17	2.16	2.15	2.15	2.14	2.14	2.13
fe 3 eq h2o ax h2o	data	2.15	2.15	2.15	2.15	2.15	2.15	2.15
	pred	2.05	2.05	2.04	2.04	2.04	2.04	2.03

fe 3 eq cl ax cl	data	2.58	2.57	2.57	2.56	2.55	2.54	2.54
fe 3 eq pisc ax cl	data	2.18	2.19	2.31	2.52	2.52	2.52	2.52
fe 3 eq c2h3ns ax c2h3ns	pred data	2.19 2.2	2.2	2.2 2.2	2.2 2.2	2.2 2.2	2.2 2.19	2.2 2.2
fe 3 eq nh3 ax scn	pred data	2.19 2.23	2.19 2.23	2.19 2.23	2.19 2.22	2.19 2.22	2.19 2.2	2.19 2.19
fe 3 eq acac ax ncs	pred data	2.24 1.92	2.23	2.22	2.22	2.21 1.92	2.21 1.93	2.2 1.93
fe 3 eq co ax scn	data	-	-	1.94	1.94	1.95 2.25	1.95 2.27	1.95 2.29
fe 3 eq bipy ax bipy	data	2.14	2.14	2.25	2.26	2.27	2.27	2.28
fe 3 eq ncs ax cn	data	2.14	2.14	2.13	2.13	2.13	2.13	2.13
fe 3 eq ncs ax ncs	data	2.12	2.12	2.12	2.12	2.12	2.09	2.08
fe 3 eq nh3 ax nh3	data	2.26	2.25	2.12	2.23	2.12	2.12	2.12
fe 3 eq cl ax pisc	data	2.16	2.17	2.18	2.19	2.18	2.19	2.19
co 2 eq ox ax ox	data	2.15	2.12	2.16	2.15	2.15	2.15	2.15
co 2 eq en ax en	data	2.13	2.13	2.13	2.13	2.13	2.13	2.13
co 2 eq tbuc ax tbuc	pred data	2.23 2.17	2.23 2.18	2.23 2.18	2.22 2.17	2.22 2.17	2.22 2.17	2.22 2.17
co 2 eq cl av pisc	pred	2.17	2.17	2.17	2.17	2.16	2.16	2.16
	pred	2.02	2.03	2.03	2.04	2.04	2.06	2.07
co z eq ch ax hzo	pred	2.02	2.01	2.02	2.03	2.04 2.04	2.05	2.06
co 2 eq en ax pisc	data pred	2.11 2.15	2.16 2.16	2.19 2.17	2.21 2.17	2.22 2.18	2.21 2.18	2.21 2.18
co 2 eq phen ax h2o	data pred	2.16 2.15	2.16 2.15	2.16 2.15	2.15 2.15	2.17 2.15	2.17 2.15	2.15 2.15
co 2 eq phen ax phen	data pred	2.15 2.17	2.16 2.16	2.16 2.16	2.18 2.15	2.18 2.15	2.18 2.15	2.18 2.14
co 2 eq h2o ax ncs	data pred	1.97 2	1.98 2	1.98 2	1.99 2	2 2	2.01 2	2.01 2.01
co 2 eq c2h3ns ax c2h3ns	data	2.18 2.18	2.19 2.18	2.2 2.18	2.2 2.19	2.21	2.22	2.22
co 2 eq nh3 ax nh3	data	2.25	2.25	2.25	2.24	2.24	2.24 2.23	2.24 2.23
co 2 eq acac ax cn	data pred	2.03	2.03	2.04 2.03	2.05 2.03	2.06 2.03	2.06 2.04	2.06 2.04
co 2 eq nh3 ax h2o	data pred	2.2 2.18	2.2 2.18	2.2 2.18	2.2	2.2 2.18	2.2 2.17	2.2 2.17
co 2 eq acac ax acac	data pred	2.07 2.08	2.08 2.08	2.08 2.08	2.08 2.08	2.08 2.08	2.1 2.08	2.1 2.09
co 2 eq ncs ax ncs	data pred	2.22 2.19	2.22 2.2	2.22 2.21	2.22 2.21	2.22 2.22	2.22 2.22	2.23 2.22
co 2 eq bipy ax bipy	data pred	2.18 2.15	2.18 2.15	2.18 2.14	2.17 2.14	1.97 2.13	-	-
co 2 eq co ax co	data pred	2.23 2.24	2.25 2.24	2.26 2.25	2.27 2.26	2.27 2.26	2.28 2.27	2.29 2.28
co 2 eq en ax h2o	data	2.17	2.17	2.17	2.17	2.16	2.16	2.15
co 2 eq h2o ax h2o	data	2.13	2.13	2.12	2.12	2.11	2.11	2.11
co 3 eq co ax nh3	data	-	2.09	2.09	2.09	2.09	2.09	2.09
co 3 eq acac ax nh3	data	2.06	2.06	2.05	2.04	2.03	2.02	2.01
co 3 eq tbuc ax pisc	data	1.98	2 02	2.03	2.08	2.11	2.02	2.02
co 3 eq nh3 ax nh3	data	2.24	2.23	2.23	2.22	2.21	2.2	2.19
co 3 eq ox ax ox	data	2.04	2.03	2.02	2.02	2 03	2 03	2 03
co 3 eq en ax en	data	2.25	2.24	2.23	2.22	2.21	2.2	2.19
co 3 eq acac ax acac	data	2.05	2.04	2.04	2.03	2.02	2.01	2 01
co 3 eq phen ax phen	data	2.16	2.16	2.18	2.19	2.19	2.19	2.19
co 3 eq cn ax cn	data	2.22	2.22	2.21	2.21	2.21	2.19	2.2
co 3 eq c2h3ns ax c2h3ns	data pred	2.15 2.16	2.16 2.16	2.16 2.16	2.16	2.16 2.16	2.16 2.16	2.16 2.16
co 3 eq co ax co	data pred	2.29 2.25	2.28 2.25	2.28 2.25	2.27 2.25	2.26 2.25	2.26 2.25	2.25 2.24
co 3 eq tbuc ax tbuc	data pred	2.09 2.09	2.09 2.09	2.08 2.09	2.07	2.1 2.08	2.11 2.08	2.11 2.08
co 3 eq h2o ax h2o	data	2.07	2.06	2.05	2.04	2.03	2.02	2.01
co 3 eq ncs ax ncs	data	2.11	2.11	2.11	2.12	2.17	2.17	2.17
co 3 eq pisc ax pisc	data	2.13	2.15	2.17	2.19	2.2	2.2	2.21
	picu	<u></u>	2.10	2.10	2.10	<u> </u>	<u> </u>	2.10

co 3 eq bipy ax bipy	data	2.14	2.14	2.18	2.19	2.18	2.18	2.18
1 1 , 1 ,	pred	2.15	2.15	2.15	2.15	2.15	2.15	2.15
co 3 eg acac ax ncs	data	1.89	1.9	1.9	1.93	1.96	1.98	2
	pred	1.92	1.92	1.93	1.93	1.94	1.94	1.95
co 3 eg ox ax ncs	data	2.05	2.04	2.02	2.02	2.02	2.02	2.02
	pred	2.04	2.04	2.04	2.03	2.02	2.02	2.01
co 3 eg acac ax co	data	2.02	2.01	2	2	1.99	1.99	1.99
•	pred	2.02	2.01	2.01	2.01	2	2	2
ni 2 eq tbuc ax tbuc	data	2.16	2.16	2.15	2.14	2.14	2.14	2.14
	pred	2.15	2.15	2.14	2.14	2.14	2.14	2.13
ni 2 eg bipy ax nh3	data	2.16	2.16	2.16	2.16	2.16	2.16	2.16
	pred	2.15	2.15	2.15	2.15	2.15	2.14	2.14
ni 2 eg en ax en	data	2.21	2.2	2.2	2.2	2.19	2.19	2.19
	pred	2.2	2.19	2.19	2.19	2.18	2.18	2.17
ni 2 eq co ax co	data	2.17	2.19	2.19	2.2	2.21	2.21	2.22
	pred	2.17	2.17	2.18	2.19	2.2	2.21	2.22
ni 2 eq nh3 ax nh3	data	2.21	2.21	2.21	2.21	2.2	2.2	2.2
	pred	2.2	2.2	2.2	2.2	2.2	2.19	2.19
ni 2 eq acac ax cn	data	2.01	2.01	2.02	2.02	2.02	2.03	2.03
	pred	2.01	2.01	2.01	2.02	2.02	2.02	2.02
ni 2 eq pisc ax pisc	data	2.11	2.12	2.13	2.14	2.14	2.15	2.16
	pred	2.09	2.09	2.1	2.11	2.11	2.12	2.13
ni 2 eg en ax h2o	data	2.13	2.13	2.13	2.13	2.13	2.12	2.12
	pred	2.14	2.14	2.14	2.14	2.14	2.14	2.14
ni 2 eq bipy ax bipy	data	2.13	2.13	2.13	2.13	2.13	2.13	2.13
	pred	2.13	2.13	2.13	2.13	2.13	2.13	2.12
ni 2 eq ncs ax ncs	data	2.17	2.17	2.17	2.17	2.17	2.17	2.18
	pred	2.17	2.17	2.17	2.17	2.17	2.17	2.17
ni 2 eq cn ax cn	data	-	-	2.34	2.34	2.34	2.35	2.35
	pred			2.33	2.33	2.34	2.34	2.34
ni 2 eq acac ax co	data	2.08	2.09	2.08	2.07	2.06	2.05	2.05
	pred	2.05	2.05	2.05	2.05	2.05	2.05	2.06
ni 2 eq c2h3ns ax c2h3ns	data	2.13	2.13	2.14	2.15	2.16	2.16	2.17
	pred	2.13	2.14	2.14	2.14	2.15	2.15	2.16
ni 2 eq acac ax nh3	data	2.1	2.1	2.1	2.1	2.1	2.09	2.09
	pred	2.08	2.08	2.08	2.08	2.08	2.08	2.08
ni 2 eq acac ax acac	data	2.05	2.05	2.07	2.07	2.06	2.06	2.06
	pred	2.06	2.06	2.06	2.05	2.05	2.05	2.05
ni 2 eq ox ax ox	data	2.13	2.13	2.12	2.12	2.11	2.11	2.11
•	pred	2.12	2.12	2.12	2.12	2.12	2.11	2.11

Table S29: RMSEs and MUEs (in Å) for minimum metal-ligand bond length prediction on test data by metal and oxidation state for both min(RLS) and min(RHS). The number of test cases is indicated in parentheses.

Species	RMS	E (Å)	MUE	Ξ (Å)
	min(RLS)	min(RHS)	min(RLS)	min(RHS)
Cr(II)	0.02 (68)	0.02 (78)	0.02	0.01
Cr(III)	0.02 (54)	0.02 (54)	0.01	0.01
Mn(II)	0.02 (65)	0.03 (49)	0.02	0.02
Mn(III)	0.02 (60)	0.05 (59)	0.01	0.03
Fe(II)	0.02 (64)	0.03 (59)	0.01	0.03
Fe(III)	0.02 (84)	0.03 (88)	0.01	0.02
Co(II)	0.03 (55)	0.04 (53)	0.02	0.03
Co(III)	0.02 (52)	0.02 (51)	0.02	0.02
Ni(II)	0.01 (36)	0.02 (47)	0.01	0.01

Table S30: molSimplify initial structure projected (g) gradients and RMS gradients with and without preliminary ANN assisted bond lengths

name		g (kcal/Å)	RMS grad. (Hartree/Bohr)
$Co(acac)_3$	default	-52	0.0487
	ANN	-5	0.0490
$\operatorname{Cr}(\operatorname{bipy})_3$	default	-48	0.0237
	ANN	29	0.0161
$Fe(acac)_3$	default	-50	0.0429
	ANN	23	0.0437
$Mn (misc)_3$	default	-57	0.0459
	ANN	-53	0.0454



Figure S28: Illustration showing definition of bond projected gradient, g, used to estimate the closeness of an initial geometry to equilibrium. The projected gradient is the scalar difference between the component of the negative energy gradient projected into the vector joining nuvlear positions of the metal (larger orange circle) and the ligand (smaller grey circle).

Table S31: Spin splitting energy predictions and data in kcal/mol, for the CSD test structures. For each complex, values from DFT using B3LYP and and ANN predictions are shown, along with the standard deviation of the ANN model

Test number	ANN	DFT/B3LYP	ANN Std. Dev.	Error
1	21.80	29.06	3.29	-7.81
2	21.90	14.20	3.28	7.70
3	-26.50	-23.90	3.25	-2.66
4	28.70	34.10	3.65	-5.35
5	32.90	35.10	3.46	-2.18
6	-17.00	-11.50	3.87	-5.53
7	12.60	35.2	4.25	-22.60
8	-13.80	-16.40	4.14	2.61
9	30.60	0.26	3.77	30.40
10	-44.50	-40.90	3.27	-3.61
11	-31.20	-15.20	3.74	-16.00
12	4.50	-23.2	5.29	27.70
13	-13.10	-16.70	4.00	3.62
14	28.30	30.70	4.60	-2.38
15	-61.00	-45.10	3.08	-15.9
16	44.40	69.70	5.06	-25.30
17	6.92	3.51	3.92	3.41
18	-19.50	-6.35	3.25	-13.10
19	-11.50	-3.14	4.27	-8.32
20	-21.90	-15.00	3.69	-6.93
21	-5.78	-1.69	3.23	-4.10
22	-11.70	-8.49	3.91	-3.19
23	-4.20	-5.36	3.52	1.16
24	6.44	2.57	3.46	3.87
25	9.44	-0.23	4.73	9.67
26	-7.91	-24.40	4.00	16.40
27	0.54	3.59	3.24	-3.04
28	-13.40	-36.90	5.56	23.50
29	-6.85	-23.90	3.30	17.10
30	-15.60	-17.20	3.76	1.63
31	-28.20	-38.30	3.26	10.10
32	-10.00	-8.98	3.60	-1.03
33	4.50	-23.40	5.29	27.90
34	-22.00	-24.00	3.29	2.01
35	-13.90	-22.40	4.03	8.55



Figure S29: Dissimilarity metrics for CSD data: errors in spin energy predictions for CSD structures are on the y-axis in kcal/mol and the minimum Tanimoto/FP2 disimilarity metric (1 - the Tanimoto index) between the CSD ligands and the training ligands is shown on the x-axis. A value of 1 inidicates no matches with the FP2 fingerprint.



Figure S30: Comparison of dissimilarity metrics for CSD data: errors in spin energy predictions for CSD structures are on the y-axis in kcal/mol and the Euclidean (left, red) and uncentered Pearson distances (gray, right) between the CSD structure and its nearest representation in dimensionless descriptor space is shown on the x-axis.

Table S32: LS bond distance predictions and data in Å, for the CSD test structures. For each complex, values from DFT using B3LYP and and ANN predictions are shown, along with the standard deviation of the ANN model

Test number	ANN	DFT/B3LYP	ANN Std. Dev.	Error
testno	nn	data	variance	error
1	2.01	1.95	0.02	0.06
2	2.01	1.90	0.02	0.11
3	1.97	2.06	0.03	-0.10
4	1.98	1.95	0.03	0.04
5	1.98	2.00	0.03	-0.02
6	1.99	1.97	0.03	0.03
7	1.97	1.96	0.02	0.01
8	2.02	2.07	0.03	-0.05
9	1.99	2.03	0.03	-0.03
10	2.02	2.03	0.03	-0.01
11	1.89	1.99	0.03	-0.10
12	2.03	2.04	0.02	-0.02
13	1.91	1.89	0.03	0.02
14	2.02	1.95	0.03	0.07
15	2.04	2.08	0.03	-0.05
16	1.99	1.91	0.03	0.09
17	2.04	2.01	0.03	0.03
18	1.94	2.00	0.02	-0.06
19	1.90	1.88	0.03	0.02
20	2.08	2.05	0.03	0.03
21	2.08	2.04	0.02	0.03
22	2.05	1.98	0.02	0.07
23	1.94	1.91	0.03	0.03
24	2.05	2.03	0.02	0.02
25	1.98	1.99	0.03	-0.01
26	2.00	1.94	0.03	0.06
27	2.04	1.97	0.02	0.06
28	2.01	2.08	0.02	-0.07
29	1.99	2.04	0.3	-0.05
30	1.88	2.04	0.03	-0.17
31	2.10	2.09	0.02	0.01
32	1.99	2.02	0.03	-0.03
33	2.03	2.04	0.02	-0.01
34	2.08	2.01	0.02	0.07
35	2.16	1.98	0.02	0.19

Table S33: HS bond distance predictions and data in Å, for the CSD test structures. For each complex, values from DFT using B3LYP and and ANN predictions are shown, along with the standard deviation of the ANN model

Test Number	ANN	DFT/B3LYP	ANN Std. Dev.	Error
1	2.06	1.95	0.02	0.11
2	2.06	1.87	0.02	0.16
3	1.94	2.07	0.02	-0.12
4	2.02	2.01	0.02	0.016
5	2.05	2.13	0.02	-0.08
6	2.02	2.06	0.02	-0.04
7	2.06	2.03	0.02	0.03
8	2.09	2.32	0.02	-0.23
9	2.00	2.05	0.02	-0.04
10	2.15	2.12	0.03	0.03
11	1.96	2.03	0.02	-0.07
12	1.98	2.06	0.02	-0.08
13	2.02	1.94	0.02	0.09
14	2.00	2.08	0.02	-0.08
15	2.21	2.22	0.02	-0.01
16	2.07	1.91	0.02	0.16
17	2.06	2.01	0.02	0.06
18	2.14	2.13	0.02	0.01
19	2.02	1.98	0.02	0.04
20	2.11	2.19	0.02	-0.08
21	2.23	2.05	0.02	0.187
22	2.09	2.07	0.02	0.02
23	2.12	2.03	0.02	0.10
24	2.24	2.16	0.02	0.08
25	1.99	2.02	0.02	-0.02
26	2.03	1.96	0.02	0.07
27	2.15	2.17	0.02	-0.02
28	2.13	2.26	0.02	-0.13
29	2.03	2.07	0.02	-0.03
30	2.12	2.08	0.02	0.04
31	2.28	2.25	0.02	0.03
32	2.19	2.22	0.02	-0.03
33	1.98	2.06	0.02	-0.09
34	2.09	2.02	0.02	0.07
35	2.21	1.99	0.02	0.22



Figure S31: Comparison of dissimilarity metrics for CSD data: errors in LS bond length prediction for CSD structures are shown on the y-axis in Å, and three normalized dissimilarity metrics are compared on the x-axis: the Tanimoto/FP2 disimilarity metric between the CSD ligands and the training ligands (blue circles), and the Euclidean (red diamonds) and uncentered Pearson distances (gray crosses) between the CSD structure and its nearest representation in dimensionless descriptor space.



Figure S32: Comparison of dissimilarity metrics for CSD data: errors in HS bond length prediction for CSD structures are shown on the y-axis in Å, and three normalized dissimilarity metrics are compared on the x-axis: the Tanimoto/FP2 disimilarity metric between the CSD ligands and the training ligands (blue circles), and the Euclidean (red diamonds) and uncentered Pearson distances (gray crosses) between the CSD structure and its nearest representation in dimensionless descriptor space.



Figure S33: Comparison of measured CSD bond distances in the crystal phase, represented by symbols (red squares for high-spin or blue triangles for low-spin based on DFT assignment at aHF=0.20) with the ANN predicted HS (red line) and LS (blue line) bond distances.



Figure S34: Parity plot comparing prediction of ΔE_{HS-LS} in kcal/mol. The x-axis is the DFT value at $a_{HF} = 0.20$, while the y-axis is the predicted value. The blue squares are obtained by interpolating from the values at $a_{HF} = 0.00$ using the predicted slopes. The red triangles are the matching test cases (where these were tested at $a_{HF} = 0.20$).
Table S34: Predicted spin splitting energies in kcal/mol for CSD structures at $a_{HF} = 0.20$ (B3LYP) interpolating data from $a_{HF} = 0$ (GGA) using ANN-predicted HFX sensitivity, compared to data from DFT using B3LYP. The predictions and standard deviations of the ANN model are given in the second and third columns in kcal/mol·HFX⁻¹.

Test	ANN $\frac{\partial E_{H-L}}{\partial a_{HF}}$	ANN Std. Dev.	$DFT,a_{\mathrm{HF}}=0.00$	Interpolated	$DFT, a_{\mathrm{HF}} = 0.20$	Error
1	-112	16.31	36.54	14.15	29.61	-15.46
2	-113.6	15.90	24.95	2.22	14.18	-11.96
3	-6.95	14.38	-22.58	-23.97	-23.87	-0.10
4	-90.68	15.10	43.97	25.83	34.05	-8.22
5	-85.80	15.01	43.98	26.82	35.05	-8.23
6	-21.31	14.56	-2.01	-6.27	-11.49	5.22
7	-101.30	14.58	37.01	16.75	35.19	-18.44
8	-126.30	13.61	3.26	-21.99	-16.41	-5.58
9	-123.10	14.90	20.23	-4.37	0.26	-4.63
11	-52.19	14.18	-8.39	-18.83	-15.21	-3.62
12	-25.51	14.53	-17.82	-22.92	-23.15	0.23
13	-59.05	14.16	-3.19	-15.00	-16.74	1.75
14	-127.00	15.08	57.46	32.07	30.66	1.41
15	-48.81	14.19	-31.03	-40.80	-45.11	4.32
16	-116.70	14.13	70.10	46.75	69.67	-22.92
17	-95.78	16.33	11.45	-7.70	3.51	-11.21
18	-46.71	13.99	5.40	-3.94	-6.35	2.41
19	-64.62	14.26	12.74	-0.18	-3.14	3.00
20	-83.78	13.97	-3.82	-20.58	-15.01	-5.56
21	-90.32	13.88	17.65	-0.42	-1.69	1.27
22	-105.20	13.67	11.70	-9.34	-8.49	-0.85
23	-80.10	14.00	8.43	-7.59	-5.36	-2.22
24	-62.35	13.85	15.41	2.94	2.57	0.37
25	-72.41	14.40	11.80	-2.69	-0.23	-2.46
26	-16.00	14.40	-21.00	-24.20	-24.36	0.15
27	-115.50	13.71	26.11	3.01	3.59	-0.58
28	-94.16	14.69	-12.56	-31.4	-36.86	5.46
29	-57.17	15.41	-17.22	-28.65	-23.90	-4.75
30	-107.30	13.88	-13.39	-34.85	-17.18	-17.67
31	-79.92	13.64	-23.14	-39.12	-38.34	-0.78
32	-69.41	14.77	2.12	-11.77	-8.98	-2.79
33	-25.51	14.53	-20.30	-25.40	-23.41	-2.00
34	-6.862	14.24	-19.72	-21.09	-23.98	2.89
35	-34.22	13.42	-18.83	-25.68	-22.43	-3.25

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