

Supplementary Materials for

Directed evolution of cytochrome c for carbon-silicon bond formation: Bringing silicon to life

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I. Materials and Methods

Unless otherwise noted, all chemicals and reagents for chemical reactions were obtained from commercial suppliers (Acros, Arch Bioscience, Fisher Scientific, Sigma-Aldrich, TCI America) and used without further purification. The following proteins were all purchased from Sigma-Aldrich: bovine serum albumin (BSA), cytochrome c(from bovine, equine heart and S. cerevisiae), peroxidase II (from horseradish), and catalase (from C. glutamicum). Silica gel chromatography purifications were carried out using AMD Silica Gel 60, 230-400 mesh. ¹H and ¹³C NMR spectra were recorded on a Bruker Prodigy 400 MHz instrument and are internally referenced to the residual solvent peak (chloroform).²⁹Si NMR spectra were recorded on the same instrument and referenced to tetramethoxysilane (δ –78.9 ppm). Data for ¹H NMR are reported in the conventional form: chemical shift (δ ppm), multiplicity (s = singlet, d = doublet, t = triplet, q = quartet, hept = heptet, m = multiplet, br = broad, app = appears as), coupling constant (Hz), integration. Data for ${}^{13}C$ and ${}^{29}Si$ are reported in terms of chemical shift (δ ppm). High-resolution mass spectra were obtained with a JEOL JMS-600H High Resolution Mass Spectrometer at the California Institute of Technology Mass Spectral Facility. Sonication was performed using a Qsonica Q500 sonicator. Chemical reactions were monitored using thin layer chromatography (Merck 60 silica gel plates) and a UVlamp for visualization. Gas chromatography (GC) analyses were carried out using a Shimadzu GC-17A gas chromatograph, a FID detector, and J&W HP-5 (30 m x 0.32 mm, 0.25 μm film; 90 °C hold 1 min, 90 to 110 °C at 15 °C/min, 110 to 280 °C at 60 °C/min, 280 °C hold 1 min, 6.2 min total). Analytical chiral supercritical fluid chromatography (SFC) was performed with a JACSO 2000 series instrument using *i*-PrOH and supercritical CO_2 as the mobile phase, with visualization at 210 nm. The following chiral columns were used: Daicel Chiralpak IC, Chiralpak AD-H, or Chiralcel OD-H (4.6 mm x 25 cm).

Plasmid pET22 was used as a cloning vector, and cloning was performed using Gibson assembly (31). The cytochrome *c* maturation plasmid pEC86 (32) was used as part of a two-plasmid system to express prokaryotic cytochrome *c* proteins. Cells were grown using Luria-Bertani medium or HyperBroth (AthenaES) with 100 µg/mL ampicillin and 20 µg/mL chloramphenicol (LB_{amp/chlor} or HB_{amp/chlor}). Cells without the pEC86 plasmid were grown with 100 µg/mL ampicillin (LB_{amp} or HB_{amp}). Primer sequences are available upon request. Electrocompetent *Escherichia coli* cells were prepared following the protocol of Sambrook *et al.* (33). T5 exonuclease, Phusion polymerase, and *Taq* ligase were purchased from New England Biolabs (NEB, Ipswich, MA). M9-N minimal medium (abbreviated as M9-N buffer; pH 7.4) was used as a buffering system for whole cells, lysates, and purified proteins, unless otherwise specified. M9-N buffer was used without a carbon source; it contains 47.7 mM Na₂HPO₄, 22.0 mM KH₂PO₄, 8.6 mM NaCl, 2.0 mM MgSO₄, and 0.1 mM CaCl₂.

II. General Procedures

(A) Plasmid construction. All variants described in this paper were cloned and expressed using the pET22(b)+ vector (Novagen). The gene encoding *Rma* cyt *c* (UNIPROT ID B3FQS5) was obtained as a single gBlock (IDT), codon-optimized for *E. coli*, and cloned using Gibson assembly (31) into pET22(b)+ (Novagen) between restriction sites *NdeI* and *XhoI* in frame with an *N*-terminal pelB leader sequence (to ensure periplasmic localization and proper maturation; MKYLLPTAAAGLLLLAAQPA MA) and a *C*-terminal 6xHis-tag. This plasmid was co-transformed with the cytochrome *c* maturation plasmid pEC86 (32) into *E. cloni*[®] EXPRESS BL21(DE3) cells (Lucigen).

(B) Cytochrome *c* expression and purification. Purified cytochrome *c* proteins were prepared as follows. One liter $HB_{amp/chlor}$ in a 4 L flask was inoculated with an overnight culture (20 mL, $LB_{amp/chlor}$) of recombinant *E. cloni*[®] EXPRESS BL21(DE3) cells containing a pET22(b)+ plasmid encoding the cytochrome *c* variant, and the pEC86 plasmid. The culture was shaken at 37 °C and 200 rpm (no humidity control) until the OD₆₀₀ was 0.7 (approximately 3 hours). The culture was placed on ice for 30 minutes, and isopropyl β -D-1-thiogalactopyranoside (IPTG) and 5-aminolevulinic acid (ALA) were added to final concentrations of 20 μ M and 200 μ M respectively.

The incubator temperature was reduced to 20 °C, and the culture was allowed to shake for 20 hours at 200 rpm. Cells were harvested by centrifugation (4 °C, 15 min, 4,000xg), and the cell pellet was stored at -20 °C until further use (at least 24 hours). The cell pellet was resuspended in buffer containing 100 mM NaCl, 20 mM imidazole, and 20 mM Tris-HCl buffer (pH 7.5 at 25 °C) and cells were lysed by sonication (2 minutes, 2 seconds on, 2 seconds off, 40% duty cycle; Qsonica Q500 sonicator). Cell lysate was placed in a 75 °C heat bath for 10 minutes, and cell debris was removed by centrifugation for 20 min (5000xg, 4 °C). Supernatant was sterile filtered through a 0.45 µm cellulose acetate filter and purified using a 1 mL Ni-NTA column (HisTrap HP, GE Healthcare, Piscataway, NJ) using an AKTA purifier FPLC system (GE healthcare). The cytochrome *c* protein was eluted from the column by running a gradient from 20 to 500 mM imidazole over 10 column volumes.

The purity of the collected cytochrome c fractions was analyzed using sodium dodecyl sulfate-polyacrylamide gel electrophoresis (SDS-PAGE). Pure fractions were pooled and concentrated using a 3 kDa molecular weight cut-off centrifugal filter and dialyzed overnight into 0.05 M phosphate buffer (pH = 7.5) using 3 kDa molecular weight cut-off dialysis tubing. The dialyzed protein was concentrated again, flash-frozen on dry ice, and stored at -20 °C.

The concentration of cytochrome c was determined in triplicate using the ferrous assay described in section (E).

(C) P450 and globin expression and purification. Purified P450s and globins were prepared differently from the cytochrome *c* proteins, and described as follows. One liter HB_{amp} in a 4 L flask was inoculated with an overnight culture (20 mL, LB_{amp}) of recombinant *E. cloni*[®] EXPRESS BL21(DE3) cells containing a pET22(b)+ plasmid encoding the P450 or globin variant. The culture was shaken at 37 °C and 200 rpm (no humidity control) until the OD₆₀₀ was 0.7 (approximately 3 hours). The culture was

placed on ice for 30 minutes, and IPTG and 5-ALA were added to final concentrations of 0.5 mM and 1 mM, respectively. The incubator temperature was reduced to 20 °C, and the culture was allowed to shake for 20 hours at 200 rpm. Cells were harvested by centrifugation (4 °C, 15 min, 4,000xg), and the cell pellet was stored at -20 °C until further use (at least 24 hours). The cell pellet was resuspended in buffer containing 100 mM NaCl, 20 mM imidazole, and 20 mM Tris-HCl buffer (pH 7.5 at 25 °C). Hemin (30 mg/mL, 0.1 M NaOH; Frontier Scientific) was added to the resuspended cells such that 1 mg of hemin was added for every 1 gram of cell pellet. Cells were lysed by sonication (2 minutes, 1 seconds on, 2 seconds off, 40% duty cycle; Qsonica Q500 sonicator). Cell debris was removed by centrifugation for 20 min (27,000xg, 4 °C). Supernatant was sterile filtered through a 0.45 µm cellulose acetate filter, and purified using a 1 mL Ni-NTA column (HisTrap HP, GE Healthcare, Piscataway, NJ) using an AKTA purifier FPLC system (GE healthcare). The P450 and globin proteins were eluted from the column by running a gradient from 20 to 500 mM imidazole over 10 column volumes.

The purity of the collected protein fractions was analyzed using SDS-PAGE. Pure fractions were pooled and concentrated using a 10 kDa molecular weight cut-off centrifugal filter and buffer-exchanged with 0.1 M phosphate buffer (pH = 8.0). The purified protein was flash-frozen on dry ice and stored at -20 °C.

P450 and globin concentrations were determined in triplicate using the hemochrome assay described in section (D).

(D) Hemochrome assay. A solution of sodium dithionite (10 mg/mL) was prepared in M9-N buffer. Separately, a solution of 1 M NaOH (0.4 mL) was mixed with pyridine (1 mL), followed by centrifugation (10,000xg, 30 seconds) to separate the excess aqueous layer gave a pyridine-NaOH solution. To a cuvette containing 700 μ L protein solution (purified protein or heat-treated lysate) in M9-N buffer, 50 μ L of dithionite solution and 250 μ L pyridine-NaOH solution were added. The cuvette was sealed with Parafilm, and the UV-Vis spectrum was recorded immediately. Cytochrome *c* concentration was determined using $\varepsilon_{550.535} = 22.1 \text{ mM}^{-1}\text{cm}^{-1}$ (34). Protein concentrations determined by the hemochrome assay were in agreement with that determined by the bicinchoninic acid (BCA) assay (Thermo Fisher) using bovine serum albumin (BSA) for standard curve preparation.

(E) Ferrous assay. To a cuvette containing 700 μ L protein solution in M9-N buffer was added 50 μ L of dithionite solution (10 mg/mL in M9-N buffer). The cuvette was sealed with Parafilm, and the UV-Vis spectrum was recorded immediately. The absorbance value for the peak at 550 nm was recorded, and background absorbance at 600 nm was subtracted. Using the protein concentration as determined by the hemochrome assay, ferrous $\varepsilon_{550-600}$ was determined to be 27 mM⁻¹cm⁻¹ for wild-type *Rma* cyt *c*, and 21 mM⁻¹cm⁻¹ for *Rma* V75T M100D M103E (see calibration curves shown on the following page). Concentrations of *Rma* M100D and V75T M100D were determined using the extinction coefficient calculated for V75T M100D M103E.



(F) Library construction. Cytochrome c site-saturation mutagenesis libraries were generated using a modified version of the 22-codon site-saturation method (35). For each site-saturation library, oligonucleotides were ordered such that the coding strand contained the degenerate codon NDT, VHG or TGG. The reverse complements of these primers were also ordered. The three forward primers were mixed together in a 12:9:1 ratio, (NDT:VHG:TGG) and the three reverse primers were mixed similarly. Two PCRs were performed, pairing the mixture of forward primers with a pET22(b)+ internal reverse primer, and the mixture of reverse primers with a pET22b internal forward primer. The two PCR products were gel purified, ligated together using Gibson assembly (31), and transformed into *E. cloni*[®] EXPRESS BL21(DE3) cells. Primer sequences are available upon request.

(G) Enzyme library screening. Single colonies were picked with toothpicks off of $LB_{amp/chlor}$ agar plates, and grown in deep-well (2 mL) 96-well plates containing $LB_{amp/chlor}$ (400 µL) at 37 °C, 250 rpm shaking, and 80% relative humidity overnight. After 16 hours, 30 µL aliquots of these overnight cultures were transferred to deep-well 96-well plates containing $HB_{amp/chlor}$ (1 mL) using a 12-channel EDP3-Plus 5-50 µL pipette (Rainin). Glycerol stocks of the libraries were prepared by mixing cells in $LB_{amp/chlor}$ (100 µL) with 50% v/v glycerol (100 µL). Glycerol stocks were stored at -78 °C in 96-well microplates. Growth plates were allowed to shake for 3 hours at 37 °C, 250 rpm shaking, and 80% relative humidity. The plates were then placed on ice for 30 min. Cultures were

induced by adding 10 μ L of a solution, prepared in sterile deionized water, containing 2 mM isopropyl β-D-1-thiogalactopyranoside (IPTG) and 20 mM ALA. The incubator temperature was reduced to 20 °C, and the induced cultures were allowed to shake for 20 hours (250 rpm, no humidity control). Cells were pelleted (4,000xg, 5 min, 4 °C) and resuspended in 500 µL M9-N buffer. For cell lysis, plates were placed in a 75 °C water bath for 10 min, followed by centrifugation (4,000xg, 5 min, 4 °C) to remove cell debris. The resulting heat-treated lysates (340 μ L) were then transferred to deep-well plates for biocatalytic reactions. In an anaerobic chamber, to deep-well plates of heat-treated lysates were added $Na_2S_2O_4$ (40 µL per well, 100 mM in dH₂O), PhMe₂SiH (10 µL per well, 400 mM in MeCN) and Me-EDA (10 µL per well, 400 mM in MeCN). The plates were sealed with aluminum sealing tape, removed from the anaerobic chamber, and shaken at 400 rpm for 1.5 h. After quenching with cyclohexane (1 mL), internal standard was added (20 µL of 20 mM methyl 2-phenylacetate in cyclohexane) and the reaction mixtures were pipetted up and down to thoroughly mix the organic and aqueous layers. The plates were centrifuged (4,000xg, 5 min) and the organic layer (400 µL) was transferred to shallowwell 96-well plates for SFC analysis. Hits from library screening were confirmed by small-scale biocatalytic reactions, which were analyzed by GC and SFC for accurate determination of turnovers and enantioselectivities.

(H) Protein lysate preparation. Protein lysates for biocatalytic reactions were prepared as follow: *E. coli* cells expressing *Rma* cyt *c* variant were pelleted (4,000xg, 5 min, 4 °C), resuspended in M9-N buffer and adjusted to the appropriate OD₆₀₀. The whole-cell solution was heat-treated (75 °C for 10 min) then centrifuged (14,000xg, 10 min, 4 °C) to remove cell debris. The supernatant was sterile filtered through a 0.45 μ m cellulose acetate filter into a 6 mL crimp vial, crimp sealed, and the head space of the crimp vial was degassed by bubbling argon through for at least 10 min. The concentration of cytochrome *c* protein lysate was determined using the ferrous assay described in section (E). Using this protocol, the protein concentrations we typically observed for OD₆₀₀ = 15 lysates are in the 8-15 μ M range for wild-type *Rma* cyt *c* and 2-10 μ M for other *Rma* cyt *c* variants.

(I) Small-scale biocatalytic reaction. In an anaerobic chamber, protein lysate (340 μ L) in a 2 mL crimp vial was added 40 μ L Na₂S₂O₄ (100 mM in dH₂O), 10 μ L PhMe₂SiH (400 or 800 mM in MeCN) and 10 μ L Me-EDA (400 mM in MeCN). The vial was crimp sealed, removed from the anaerobic chamber, and shaken at 400 rpm at room temperature for the stated reaction time. At the end of the reaction, the crimp vial was opened and the reaction was quenched with cyclohexane (1 mL). Internal standard was added (20 μ L of 20 mM 2-phenylethanol in cyclohexane) and the reaction mixture was transferred to a microcentrifuge tube, vortexed (10 seconds, 3 times), then centrifuged (14,000xg, 5 min) to completely separate the organic and aqueous layers (the vortex-centrifugation step was repeated if complete phase separation was not achieved). The organic layer (750 μ L) was removed for GC and SFC analysis. All biocatalytic reactions were performed in triplicate unless otherwise stated. The total turnover numbers (TTNs) reported are calculated with respect to the protein catalyst and represent the total number of turnovers that is possible to obtain from the catalyst under the stated reaction conditions.

III. Supporting Tables and Figures

Chiral catalytic system	Ref	Reaction condition	Substrate scope	TTN	% ee
Copper					
Α	(15)	CH ₂ Cl ₂ , rt, 18 h	4	17 to 25	17 to 98
В	(36)	C ₆ H ₆ , 0 °C, 13.5 h	1	8 to 10	29 to 78
С	(14)	CH₂Cl₂, −60 to 0 °C, 2-12 h	24	3 to 19	12 to 99
В	(37)	CH ₂ Cl ₂ , −40 to 0 °C, 48-72 h	8	5 to 9	49 to 88
В	(38)	CH ₂ Cl ₂ , −40 to −10 °C	9	5 to 8	40 to 84
Iridium					
D	(13)	CH ₂ Cl ₂ , −78 or −30 °C, 24 h	15	24 to 50	94 to 99
E	(39)	CH₂Cl₂, −78 °C, 24 h	7	75 to 94	72 to 91
Rhodium					
F	(12)	CH ₂ Cl ₂ , rt to 40 °C, 6-12 h	34	9 to 30	77 to 99
G	(40)	CH ₂ Cl ₂ , −78 °C or rt, 3-12 h	5	8 to 45	20 to 63
н	(36)	C₅H ₁₂ , −78 to −75 °C, 24 h	1	24 to 70	48 to 97
1	(11)	CF₃CH₂OH, −35 °C	10	51 to 97	20 to 99
н	(41)	CH ₂ Cl ₂ , −78 °C then rt, 23 h	6	22 to 54	77 to 94
various Rh(II)- carboxylate	(42) (43)	CH₂Cl₂, rt or −78 °C, 0.5 to 24 h to overnight	6	<1 to 23	6 to 76
J	(44)	CH₂Cl₂, −78 °C to rt, 23 h	2	35 to 40	38 to 58
н	(45)	CH₂Cl₂, −78 °C, 1 to 1.5 h	4	34 to 43	35 to 72
н	(46)	C₅H ₁₂ , −78 °C, 24 h	5	13 to 19	75 to 95
various Rh(II)- carboxylate	(47)	CH ₂ Cl ₂ , rt or reflux	1	8 to 17	6 to 47

Table S1. Summary of known catalytic systems for enantioselective carbene insertion

 into silicon-hydrogen bonds

Catalytic systems that could yield enantiopure products are highlighted in blue. rt = room temperature. In cases where the reaction times were not documented in the original literature, this information is not shown in the table above.



Table S2. Summary of known catalytic systems that accept α -alkyl diazo compounds as substrates for enantioselective carbene insertion into silicon–hydrogen bonds

 α -Alkyl diazo compounds are challenging substrates for intermolecular carbene-transfer chemistry due to their propensity to undergo competing intramolecular β -hydride migration (48, 13). As a result, only a subset of catalytic systems shown in Table S1 have been reported to accommodate these substrates, as summarized in the table below:

Chiral catalytic system	Ref	Reaction condition	Substrate scope	TTN	% ee
Copper					
Α	(15)	CH ₂ Cl ₂ , rt, 18 h	4	17 to 25	17 to 98
С	(14)	CH₂Cl₂, −40 °C, 2-12 h	2	3 to 12	12 to 35
Iridium					
D	(13)	CH₂Cl₂, −78 °C, 24 h	8	24 to 44	94 to 99
Rhodium					
F	(12)	CH ₂ Cl ₂ , rt to 40 °C, 6-12 h	2	11 to 15	70 to 77
I	(11)	CF₃CH₂OH, −35 °C	1	n.d. [§]	64
н	(41)	CH_2CI_2 , –78 °C then rt, 23 h	5	22 to 54	77 to 94

See Table S1 for chemical structures of chiral catalysts. rt = room temperature. [§] The amount of product formed was not reported in the original literature.

Table S3. Summary of known catalytic systems for reaction between phenyldimethylsilane and Me-EDA *via* enantioselective carbene insertion into silicon–hydrogen bond



Chiral catalytic system	Ref	Reaction condition	TTN	% ee
Copper				
С	(14)	CH₂Cl₂, −40 °C, 2-12 h	12	35
Iridium				
D	(13)	CH₂Cl₂, −78 °C, 24 h	43	97
Rhodium				
F	(12)	CH ₂ Cl ₂ , rt to 40 °C, 6-12 h	15	77

See Table S1 for chemical structures of chiral catalysts. rt = room temperature.

Table S4. Preliminary experiments with heme and purified heme proteins

O L	Catalyst	•	_	
	M9-N buffer (pH 7.4 room temperature,), Na ₂ S ₂ O ₄ 1.5 hours		Si、
Catalyst		TTN	% ee	
Controls				
None		0	-	
Hemin		4 ± 1	0	
Hemin + BSA		1 ± 1	0	
P450s				
BM3 P450 T268	3A (22)	44 ± 15	0	
BM3 P450 T268	3A C400H (<i>49</i>)	45 ± 3	<5	
BM3 P450 CIS	I263F C400S T438S (50)	24 ± 9	<5	
BM3 P450 F87	A T268A C400S (<i>51</i>)	40 ± 15	<5	
BM3 P450 Hsta	r H92N H100N (52)	46 ± 7	0	
Myoglobins				
Sperm whale M	b	12 ± 1	0	
Sperm whale M	b H64V V68A (53)	17 ± 1	0	
Cytochromes	С			
Horse heart cyt	С	31 ± 10	<5	
Bovine heart cy	t c	54 ± 2	6	
S. cerevisiae cy	rt c	11 ± 1	<5	
<i>R. marinus</i> cyt	C	34 ± 10	97	
H. thermophilus	s cyt c	8 ± 2	16	
R. globiformis c	yt c	4 ± 1	<5	
Others				
Horse radish pe	eroxidase	0	-	
C. glutamicum	catalase	0	-	

Heme proteins that are available commercially or in our laboratory inventory were screened to identify the most enantioselective protein variant as starting point for directed evolution. Experiments with heme proteins were performed using 10 μ M purified heme protein, 10 mM silane, 10 mM diazo ester, 10 mM Na₂S₂O₄, 5 vol% MeCN, M9-N buffer at room temperature under anaerobic conditions for 1.5 h. Experiments with hemin were performed using 100 μ M hemin. Experiments with hemin and BSA were performed using 100 μ M hemin in the presence of BSA (0.75 mg/mL). Reactions were performed in triplicate. TTNs reported are the average of three experiments. Within instrument detection limit, variability in % *ee* was not observed. Unreacted starting materials were observed at the end of all reactions and no attempt was made to optimize these reactions.

Table S5. Carbon-silicon bond formation ca	atalyzed by <i>Rma</i> cyt c	variants
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Experiments were performed using lysates of *E. coli* expressing *Rma* cyt *c* variant (OD_{600} = 15; heat-treated at 75 °C for 10 min), 10 mM silane, 10 mM diazo ester, 10 mM Na₂S₂O₄, 5 vol% MeCN, M9-N buffer at room temperature under anaerobic conditions for 1.5 h. Reactions were performed in triplicate. TTNs reported are the average of three experiments.

Table S6. Comparison of carbon–silicon bond forming rates of four generations of *Rma* cyt *c*



To determine the initial reaction rate, experiments were performed using purified *Rma* cyt *c* variant (0.8 μ M), 10 mM silane, 10 mM diazo ester, 10 mM Na₂S₂O₄, 5 vol% MeCN, M9-N buffer at room temperature under anaerobic conditions for various time intervals. See below for experimental procedure. The data used for determining the initial rate of each *Rma* cyt *c* variant is showed in the following graph:



<i>Rma</i> cyt <i>c</i> variant	initial rate / µM min⁻¹	turnover frequency (TOF) / min ⁻¹	TOF relative to WT
WT	5.1 ± 0.3	6.4 ± 0.3	1
M100D	14.0 ± 0.4	17.5 ± 0.5	2.8 ± 0.2
V75T M100D	23.6 ± 0.8	29.5 ± 1.0	4.6 ± 0.3
V75T M100D M103E	36.4 ± 0.5	45.5 ± 0.7	7.1 ± 0.4

Errors quoted in the table above are calculated from the standard deviations of the fitting of data in the product *vs* time plot.

General procedure for carrying out timed experiments

In an anaerobic chamber, 1 mL Na₂S₂O₄ (100 mM in dH₂O) was added to 4 mL purified *Rma* cyt *c* protein (2.0 μ M in M9-buffer) to give **Solution 1**. To four 2 mL microcentrifuge tubes were each added 180 μ L M9-N buffer, 10 μ L PhMe₂SiH (400 mM in MeCN) and 10 μ L Me-EDA (400 mM in MeCN), and the mixtures were mixed

thoroughly on a shaker (480 rpm for 2 min). To these mixtures were added 200 μ L **Solution 1**, and the microcentrifuge tubes were closed and quickly shaken by hand for 3 seconds to ensure thorough mixing, before the tubes were returned to the shaker. The reactions were stopped at specific time points (2 min, 4 min, 6 min, and 8 min) by quick addition (within 10 seconds) of 40 μ L pyridine solution (400 mM in dH₂O), 20 μ L internal standard (20 mM acetophenone in toluene) and 1 mL cyclohexane. (Note: pyridine was added as a quencher to significantly slow down the reaction.) After the mixtures were vortexed for 20 seconds, 200 μ L organic layer was immediately removed for GC analysis.

Table S7. Rh(II) or Cu(II)-catalyzed reactions between Me-EDA and 4-(dimethylsilyl)phenol (1k) or 4-(dimethylsilyl)aniline (23)

 $Rh_2(OAc)_4$ and $Cu(OTf)_2$, which are known to catalyze carbene insertion into Si-H bonds under ligand-free conditions (54), were tested for their chemoselectivities towards Si-H, O-H and N-H insertions:



	21 (Si-H insertion)	21 - <i>iso</i> (O-H insertion)	21- <i>di</i> (double insertion)
Rh ₂ (OAc) ₄	×	\checkmark	\checkmark
Cu(OTf) ₂	×	×	×
	22 (Si-H insertion)	22-<i>iso</i> (N-H insertion)	22- <i>di</i> (double insertion)
Rh ₂ (OAc) ₄	×	\checkmark	×
Cu(OTf) ₂	×	×	×

Note: \times = not detected; \checkmark = detected. Cu(OTf)₂ gave complex mixtures of products in both reactions.

Experimental procedure

To a 5 mL vial was added silane (0.1 mmol, 1.0 equiv.), metal catalyst $(Rh_2(OAc)_4 (0.44 \text{ mg}, 1 \text{ mol}\%) \text{ or } Cu(OTf)_2 (3.62 \text{ mg}, 10 \text{ mol}\%))$ and DCM (0.5 mL). The mixture was cooled to -78 °C before a solution of Me-EDA (25 µL, 2.0 equiv.) in DCM (0.3 mL) was added dropwise. After slowly warming up to room temperature in 4 hours, the reaction mixture was filtrated through a short pad of silica, diluted with DCM, and analyzed by GC-MS.

Products isolated from $Rh_2(OAc)_4$ -catalyzed reactions with **1k** and **23** are shown on the next page. Both reactions generated multiple products, rendering product purification and quantitative analysis of these reactions difficult. Multiple rounds of purification by silica column chromatography were required to obtain samples suitable for characterization; yields were therefore not determined for these reactions. Notably, Si–H insertion products **21** and **22** were not observed in these reactions. (Table S7 continued)

 $Rh_2(OAc)_4$ -catalyzed reaction with 1k gave *rac*-21-*di* and *rac*-21-*iso*:





¹H NMR (400 MHz, CDCl₃) δ 7.40 (d, *J* = 8.6 Hz, 2H), 6.87 (d, *J* = 8.7 Hz, 2H), 4.75 (q, *J* = 6.8 Hz, 1H), 4.22 (q, *J* = 7.1 Hz, 2H), 4.01 (q, *J* = 7.1 Hz, 2H), 2.21 (q, *J* = 7.1 Hz, 1H), 1.62 (d, *J* = 6.8 Hz, 3H), 1.25 (t, *J* = 7.1 Hz, 3H), 1.17-1.10 (m, 6H), 0.33 (s, 6H). ¹³C NMR (100 MHz, CDCl₃) δ 176.19, 172.25, 158.86, 135.54, 128.29, 128.28, 114.65, 72.42, 61.46, 59.93, 30.30, 18.68, 14.47, 14.28, 11.43, -3.77, -4.60. HRMS (FAB) *m*/*z*: 352.1714 (M⁺); calc. for C₁₈H₂₈SiO₅: 352.1706.





¹H NMR (400 MHz, CDCl₃) δ 7.44 (d, J = 8.7 Hz, 2H), 6.87 (d, J = 8.7 Hz, 2H), 4.76 (q, J = 6.8 Hz, 1H), 4.39 (hept, J = 3.7 Hz, 1H), 4.22 (q, J = 6.9 Hz, 2H), 1.62 (d, J = 6.8 Hz, 3H), 1.26 (t, J = 7.1 Hz, 3H), 0.31 (d, J = 3.8 Hz, 6H).

 $Rh_2(OAc)_4$ -catalyzed reaction with 23 gave *rac-22-iso* and the corresponding double N–H insertion product:



¹H NMR (400 MHz, CDCl₃) δ 7.24-7.17 (m, 2H), 6.88-6.68 (m, 3H), 4.68 (dq, *J* = 5.5, 2.8 Hz, 1H), 4.22-4.11 (m, 3H), 1.50 (br d, *J* = 6.4 Hz, 3H), 1.25 (t, *J* = 7.1 Hz, 3H), 0.19 (d, *J* = 2.8 Hz, 6H). ¹³C NMR (100 MHz, CDCl₃) δ 173.94, 135.20, 129.50, 119.78, 114.93, 61.47, 53.20, 18.62, 14.31, 0.71. HRMS (FAB) *m/z*: 250.1253 ((M+H)–H₂⁺); calc. for C₁₃H₂₀SiNO₂: 250.1263.

Interestingly, the following compound was also isolated as a mixture of diastereomers. The counter anion of the ammonium salt was not determined.



¹H NMR (400 MHz, CDCl₃) δ 7.22 (overlapping doublets, J = 8.8, 7.3 Hz, 2H), 6.84 (t, J = 7.3 Hz, 1H), 6.78 (br d, J = 7.9 Hz, 2H), 4.99-4.55 (m, 1H), 4.44 (q, J = 7.1 Hz, 2H), 4.23 (app qd, J = 7.2, 2.0 Hz, 4H), 1.55 (d, J = 7.2 Hz, 6H), 1.29 (t, J = 7.1 Hz, 6H), 0.30-0.19 (m, 6H). ¹³C NMR (100 MHz, CDCl₃) δ 174.40, 146.38, 128.89, 119.30, 117.17, 61.24, 56.24, 15.95, 14.34, 0.71. HRMS (FAB) m/z: 352.1958 (M⁺); calc. for C₁₈H₃₀SiNO₄: 352.1944. Fig. S1 Representative SDS-PAGE gel of purified wild-type *Rma* cyt *c* and its V75T M100D M103E variant (TDE) in comparison to a standard protein ladder. The second and third lanes from the left are the same samples loaded at lower protein concentration.



Fig. S2 Circular dichroism (CD) spectra of purified *Rma* cyt *c* V75T M100D M103E (Rma TDE).

Rma cyt *c* V75T M100D M103E was purified without performing the heat treatment step. Two identical samples were prepared in M9-N buffer, and one was heat treated at 75 °C for 10 minutes. After cooling to room temperature, the samples were analyzed by CD. The CD spectra of heat-treated and untreated *Rma* cyt *c* V75T M100D M103E are identical, suggesting that heat treatment at 75 °C for 10 minutes does not cause irreversible denaturation of the protein. The $\Delta \varepsilon_{MRW}$ values shown here are similar to previously published values for wild-type *Rma* cyt *c* (26), which suggests mutations V75T, M100D, and M103E are not highly disruptive to the protein secondary structure.







(A) Putative active site structure of wild-type Rma cyt c showing residues V75, M100, and M103 (PDB ID: 3CP5). (B) Proposed binding mode for the iron-carbenoid, where the carbenoid forms in a way that takes the place of the axial methionine. The silane is proposed to approach from the more solvent-exposed side in the wild-type protein, which explains the observed stereochemistry of the organosilicon product. The V75T, M100D, and M103E mutations may promote reactivity by improving solvent and substrate access to the iron center. It should be noted that the working hypothesis proposed here is highly speculative as structural data about the ability of bacterial cytochrome c proteins to bind organic molecules is practically non-existent.

Fig. S4 Example of reaction time course of purified *Rma* cyt *c* M100D V75T M103Ecatalyzed reaction between phenyldimethylsilane and Me-EDA.



Experiments were performed using 3 μ M purified *Rma* cyt *c* V75T M100D M103E, 10 mM silane, 10 mM diazo ester, 10 mM Na₂S₂O₄, 5 vol% MeCN, M9-N buffer at room temperature under anaerobic conditions. Reactions were performed in duplicate. TTNs shown are the average of two experiments.

It is possible that catalysis slows down and ultimately stops due to catalyst inactivation by carbene transfer to the protein, an inactivation mechanism we recently analyzed in detail in another carbene-transfer enzyme (52). The activity and lifetime of Rma cyt cwould likely improve by identifying the protein residues affected by mechanism-based inhibitors and further mutagenesis studies. Furthermore, inactivation of whole-cell catalyst was previously found to be slower than purified enzyme (52); this might also be applicable to Rma cyt c in whole bacterial cell. Fig. S5 Additional substrates tested for *Rma* cyt *c* V75T M100D M103E-catalyzed carbon–silicon bond formation.



Rma cyt *c* V75T M100D M103E-catalyzed reactions of silanes **Si-A-H** were tested with Me-EDA and that of diazo compounds **diazo-A-C** were tested with phenyldimethylsilane under standard *in vitro* biocatalytic reaction conditions. Product formation was analyzed by GC-MS only, and comparison with chemically synthesized reference standards was not made. These preliminary results should be interpreted with caution and should not be used alone for drawing conclusions.

Si-A-D and diazo-A: formation of organosilicon product was detected by GC-MS.

Si-E-H and diazo-B-C: formation of organosilicon product was not detected by GC-MS.

Because biocatalysts that are fully genetically encoded can be readily optimized by directed evolution, the *Rma* cyt c V75T M100D M103E variant presented in this report is not the end point, but rather a parent to hundreds and thousands of future enzyme variants, some of which will most certainly surpass the scope and activity of the parent enzyme. For substrate scope expansion or other future applications, we suggest using the substrate of interest for directed evolution, and carrying out reaction screening under condition that suits individual substrate.

Fig. S6 Hammett analysis of *Rma* cyt c V75T M100D M103E-catalyzed carbon–silicon bond formation



The Hammett plot suggests a small build-up of positive charge on the silane in the reaction transition state. This observation is similar to that reported for carbene insertion into Si–H bond catalyzed by copper ($\rho = -0.54$) (37) and rhodium ($\rho = -0.31$) (55).

-0.32

0.45

0.47

CO₂Me

IV. Substrate Synthesis and Characterization

Commercially available substrates were used as received: phenyldimethylsilane (Sigma-Aldrich), benzyldimethylsilane (Sigma-Aldrich), ethyl 2-diazopropanoate (Arch Bioscience). The following diazo compounds are known and prepared according to literature procedures: isopropyl 2-diazopropanoate (56), ethyl 2-diazobutanoate (57).

Dimethyl(*p*-tolyl)silane (1b)



MeO

In a 100 mL round-bottom flask, chlorodimethylsilane (1.11 mL, 10.0 mmol) in THF (6 mL) was cooled to 0 °C. A solution of 4methylphenylmagnesium bromide (24 mL, 0.5 M in THF) was added dropwise slowly over 15 min. Then reaction was allowed to warm to room temperature and stirred for 8 hours. The reaction mixture was quenched with NH_4Cl (5 mL, sat. aq.) and the product was extracted with Et_2O (15 mL × 3). The organic layer was washed with water (20 mL), brine (20 mL), dried over MgSO₄, and then concentrated under reduced pressure (200 torr). The crude product was purified by silica column chromatography with pentane to afford **1b** (1.27 g, 8.44 mmol, 84%). This compound is known (58). ¹H NMR (400 MHz, CDCl₂) δ 7.48 (d, J = 7.8 Hz, 2H), 7.23 (d, J = 7.4 Hz, 2H), 4.46 (hept, J = 3.7 Hz, 1H), 2.39 (s, 3H), 0.37 (d, J = 3.7 Hz, 6H).

(4-Methoxyphenyl)dimethylsilane (1c)



warm to room temperature and stirred for 8 hours. The reaction mixture was quenched with NH₄Cl (5 mL, sat. aq.) and the product was extracted with Et₂O (15 mL \times 3). The organic layer was washed with water (20 mL), brine (20 mL), dried over MgSO₄, and then concentrated under reduced pressure (200 torr). The crude product was purified by silica column chromatography with pentane / Et_2O (10: 1) to afford 1c (1.60 g, 9.62 mmol, 96%). This compound is known (58). ¹H NMR (400 MHz, CDCl₂) δ 7.51 (d, J = 8.7 Hz, 2H), 6.96 (d, J = 8.6 Hz, 2H), 4.75-4.19 (m, 1H), 3.85 (s, 3H), 0.37 (d, J = 3.8Hz, 6H).

(4-Chlorophenyl)dimethylsilane (1d)



In a 100 mL round-bottom flask, chlorodimethylsilane (1.11 mL, 10.0 mmol) in THF (6 mL) was cooled to 0 °C. A solution of 4chlorophenylmagnesium bromide (24 mL, 0.5 M in THF) was added dropwise slowly over 15 min. Then reaction was allowed to warm to

room temperature and stirred for 8 hours. The reaction mixture was quenched with

NH₄Cl (5 mL, sat. aq.) and the product was extracted with Et₂O (15 mL × 3). The organic layer was washed with water (20 mL), brine (20 mL), dried over MgSO₄, and then concentrated under reduced pressure (200 torr). The crude product was purified by silica column chromatography with pentane to afford **1d** (1.27 g, 7.45 mmol, 75%). This compound is known (58). ¹H NMR (400 MHz, CDCl₃) δ 7.46 (d, *J* = 8.3 Hz, 2H), 7.34 (d, *J* = 8.2 Hz, 2H), 4.41 (hept, *J* = 3.8 Hz, 1H), 0.34 (d, *J* = 3.7 Hz, 6H).

(4-(Trifluoromethyl)phenyl)dimethylsilane (1e)

In a 100 mL round-bottom flask, 1-bromo-4-(trifluoromethyl)benzene (1.4 mL, 10.0 mmol) in THF (15 mL) was cooled to -78 °C. *n*-BuLi (7.5 mL, 1.6 M in hexane) was added dropwise slowly over 15 min. The resulting mixture was stirred at -78 °C for 2 hours before the

dropwise addition of chlorodimethylsilane (1.0 mL, 9.0 mmol). The reaction was allowed to warm to room temperature and stirred for 8 hours. The reaction mixture was quenched with NH₄Cl (5 mL, sat. aq.) and the product was extracted with Et₂O (15 mL × 3). The organic layer was washed with water (20 mL), brine (20 mL), dried over MgSO₄, and then concentrated under reduced pressure (200 torr). The crude product was purified by silica column chromatography with pentane to afford **1e** (0.81 g, 3.97 mmol, 40%). This compound is known (58). ¹H NMR (400 MHz, CDCl₃) δ 7.66 (d, *J* = 7.7 Hz, 2H), 7.60 (d, *J* = 7.9 Hz, 2H), 4.46 (hept, *J* = 3.8 Hz, 1H), 0.38 (d, *J* = 3.8 Hz, 6H).

(4-(Chloromethyl)phenyl)dimethylsilane (1f)



In a 250 mL round-bottom flask, (4-bromophenyl)methanol (5.61 g, 30.0 mmol) in THF (100 mL) was cooled to -78 °C. *n*-BuLi (30.0 mL, 2.5 M in hexane) was added dropwise slowly over 30 min. The resulting mixture was stirred at -78 °C for 2 hours before the

dropwise addition of chlorodimethylsilane (4.5 mL, 40.0 mmol). The reaction was allowed to warm to room temperature and stirred for 8 hours. The reaction mixture was quenched with NH₄Cl (50 mL, sat. aq.) and the product was extracted with DCM (50 mL × 3). The organic layer was washed with water (20 mL), brine (20 mL), dried over MgSO₄, and then concentrated under reduced pressure (100 torr). The crude product was purified by silica column chromatography with EtOAc / hexane (1:3) to afford (4-(dimethylsilyl)phenyl)methanol (2.96 g, 17.8 mmol, 59%). This compound is known (59). ¹H NMR (400 MHz, CDCl₃) δ 7.55 (d, *J* = 8.0 Hz, 2H), 7.37 (d, *J* = 8.1 Hz, 2H), 4.70 (s, 2H), 4.43 (hept, *J* = 3.7 Hz, 1H), 0.35 (d, *J* = 3.8 Hz, 6H).

To a solution of (4-(dimethylsilyl)phenyl)methanol (498.9 mg, 3.0 mmol) in DCM (4 mL) were added triethylamine (0.5 mL, 3.6 mmol) and 4-methylbenzenesulfonyl chloride (629.1 mg, 3.3 mmol). The reaction mixture was stirred at room temperature for 8 hours. The reaction was then diluted with DCM (10 mL) and washed with water (20 mL), brine (20 mL), dried over MgSO₄, and the organic layer was concentrated under reduced pressure (100 torr). The crude product was purified by silica column chromatography with pentane to afford **1f** (0.22 g, 1.19 mmol, 40%). ¹H NMR (400

MHz, CDCl₃) δ 7.55 (d, J = 8.0 Hz, 2H), 7.39 (d, J = 7.9 Hz, 2H), 4.59 (s, 2H), 4.43 (hept, J = 3.8 Hz, 1H), 0.35 (d, J = 3.8 Hz, 6H). HRMS (FAB) m/z: 183.0399 ((M+H)– H_2^+); calc. for C₉H₁₂SiCl: 183.0397.

Methyl 4-(dimethylsilyl)benzoate (1g)



In a 100 mL round-bottom flask, methyl 4-iodobenzoate (2.62 g, 10.0 mmol) in THF (15 mL) was cooled to -78 °C. *i*-PrMgCl (6 mL, 2.0 M in Et₂O) was added dropwise slowly over 5 min. The resulting mixture was allowed to warm to -40 °C in 2 hours and maintained at -40 °C for another 2 hours before the dropwise

addition of chlorodimethylsilane (1.2 mL, 11.0 mmol). The reaction was allowed to warm to room temperature and stirred for 8 hours. The reaction mixture was quenched with NH₄Cl (5 mL, sat. aq.) and the product was extracted with DCM (15 mL × 3). The organic layer was washed with water (20 mL), brine (20 mL), dried over MgSO₄, and then concentrated under reduced pressure (100 torr). The mixture was dissolved in Et₂O (5 mL) and treated with hexane (25 mL). This crashed out most of the starting material, which was removed by filtration. The filtrate was collected, concentrated under reduced pressure, and purified by silica column chromatography with EtOAc / hexane (1:20) to afford **1g** (0.88 g, 4.53 mmol, 45%). ¹H NMR (400 MHz, CDCl₃) δ 8.00 (d, *J* = 8.1 Hz, 2H), 7.62 (d, *J* = 8.2 Hz, 2H), 4.45 (hept, *J* = 3.8 Hz, 1H), 3.92 (s, 3H), 0.37 (d, *J* = 3.8 Hz, 6H). ¹³C NMR (100 MHz, CDCl₃) δ 167.35, 143.92, 134.13, 130.78, 128.72, 52.27, – 3.84. HRMS (FAB) *m/z*: 195.0843 (M+H⁺); calc. for C₁₀H₁₅SiO₂: 195.0841.

4-(dimethylsilyl)-*N*,*N*-dimethylbenzamide (1h)



To a solution of dimethylamine hydrochloride (2.45 g, 30.0 mmol) in DCM (50 mL) was added triethylamine (4.2 mL, 30.0 mmol). The mixture was stirred for 30 min before the addition of 4-iodobenzoic acid (6.20 g, 25.0 mmol) and 1-(3-dimethylaminopropyl)-3-ethylcarbodiimide (EDC, 4.66 g, 30.0

mmol). The reaction was stirred for 8 hours at room temperature. Then the reaction mixture was washed with water (50 mL), HCl (aq., 1 M, 50 mL), NaHCO₃ (sat. aq., 50 mL) and brine (50 mL), dried over MgSO₄, and then concentrated under reduced pressure (50 torr). The crude product was dissolved in Et_2O (5 mL) and treated with hexane (50 mL). The target product 4-iodo-*N*,*N*-dimethylbenzamide (6.88 g, 25.0 mmol, quantitative) crashed out and was collected by filtration.

In a 100 mL round-bottom flask, 4-iodo-*N*,*N*-dimethylbenzamide (1.65 g, 6.0 mmol) in THF (15 mL) was cooled to -78 °C. *i*-PrMgCl (6 mL, 2.0 M in Et₂O) was added dropwise slowly over 5 min. The resulting mixture was allowed to warm to -40 °C within 2 hours and maintained at -40 °C for another 2 hours before the dropwise addition of chlorodimethylsilane (1.3 mL, 12.0 mmol). The reaction was allowed to warm to room temperature and stirred for 8 hours. The reaction mixture was quenched with NH₄Cl (5 mL, sat. aq.) and the product was extracted with DCM (15 mL × 3). The organic layer

was washed with water (20 mL), brine (20 mL), dried over MgSO₄, and then concentrated under reduced pressure (50 torr). The mixture was dissolved in Et₂O (5 mL) and treated with hexane (25 mL). Most of the starting material crashed out and was removed by filtration. The filtrate was collected, concentrated under reduced pressure, and then purified by silica column chromatography with EtOAc / hexane (1:2) to afford **1h** (0.12 g, 0.579 mmol, 10%). ¹H NMR (400 MHz, CDCl₃) δ 7.57 (d, *J* = 8.3 Hz, 2H), 7.39 (d, *J* = 8.3 Hz, 2H), 4.43 (hept, *J* = 3.8 Hz, 1H), 3.05 (s, 6H), 0.35 (d, *J* = 3.7 Hz, 6H). ¹³C NMR (100 MHz, CDCl₃) δ 171.74, 139.53, 137.04, 134.11, 126.42, -3.74. HRMS (FAB) *m/z*: 208.1153 (M+H⁺); calc. for C₁₁H₁₈ONSi: 208.1158.

(3,4-Dihydro-2*H*-pyran-6-yl)dimethylsilane (1i)

In a 100 mL round-bottom flask, 3,4-dihydro-2*H*-pyran (2.00 g, 24.0 mmol) in THF (1.0 mL) and pentane (40 mL) was cooled to -78 °C. *t*-BuLi (15.5 mL, 1.7 M in pentane) was added dropwise slowly over 20 min. The resulting mixture was allowed to warm to 0 °C within 2 hours and maintained at 0 °C for another 2 hours before the dropwise addition of chlorodimethylsilane (2.6 mL, 24.0 mmol). The reaction was allowed to warm to room temperature and stirred for 8 hours. The reaction mixture was quenched with NH₄Cl (5 mL, sat. aq.) and the product was extracted with Et₂O (15 mL × 3). The organic layer was washed with water (20 mL), brine (20 mL), dried over MgSO₄, and then concentrated under reduced pressure (200 torr). The crude product was purified by silica column chromatography with Et₂O / pentane (1: 30) to afford **1i** (2.80 g, 19.7 mmol, 82%). This compound is known (60). ¹H NMR (400 MHz, CDCl₃) δ 5.09 (t, J = 3.8 Hz, 1H), 4.05-3.91 (m, 3H), 2.04 (td, J = 6.4, 3.8 Hz, 2H), 1.92-1.83 (m, 2H), 0.19 (d, J = 3.8 Hz, 6H).

4-(Dimethylsilyl)phenol (1k)



To a solution of 4-bromophenol (1.73 g, 10.0 mmol) in THF (15 mL) was added NaH (60% in mineral oil, 0.48 g, 12.0 mmol). After the mixture was stirred for 30 min, it was cooled down to -78 °C. *t*-BuLi (6 mL, 1.7 M in pentane) was added dropwise slowly over 15 min.

The resulting mixture was stirred at -78 °C for 2 hours before the dropwise addition of chlorodimethylsilane (2.3 mL, 21.0 mmol). The reaction was allowed to warm to room temperature and stirred for 8 hours. The reaction mixture was quenched with NH₄Cl (5 mL, sat. aq.) and the product was extracted with DCM (15 mL × 3). The organic layer was washed with water (20 mL), brine (20 mL), dried over MgSO₄, and then concentrated under reduced pressure (50 torr). The crude product was purified by silica column chromatography with EtOAc / hexane (1:7) to afford **1k** (0.60 g, 3.94 mmol, 39%). This compound is known (*61*). ¹H NMR (400 MHz, CDCl₃) δ 7.43 (d, *J* = 8.5 Hz, 2H), 6.85 (d, *J* = 8.5 Hz, 2H), 4.78 (s, 1H), 4.40 (hept, *J* = 3.7 Hz, 1H), 0.32 (d, *J* = 3.8 Hz, 6H). ¹³C NMR (100 MHz, CDCl₃) δ 156.61, 135.82, 128.66, 115.16, -3.38.

(4-(Benzyloxy)phenyl)dimethylsilane (1ks)



To a solution of 4-bromophenol (5.19 g, 30.0 mmol) in MeCN (50 mL) was added K_2CO_3 (5.53 g, 40.0 mmol). After the mixture was stirred at 40 °C for 30 min, BnBr (3.6 mL, 30.0 mmol) was added over 2 min. The resulting mixture was stirred at 40 °C for 3 hours. The reaction mixture was washed

with water (20 mL), brine (20 mL), dried over $MgSO_4$, and then concentrated under reduced pressure (50 torr) to afford the crude product 1-(benzyloxy)-4-bromobenzene (7.85 g, 30.0 mmol, quantitative).

In a 100 mL round-bottom flask, 1-(benzyloxy)-4-bromobenzene (6.57 g, 25.0 mmol) in THF (15 mL) was cooled to -78 °C. *t*-BuLi (19.0 mL, 1.7 M in pentane) was added dropwise slowly over 15 min. The resulting mixture was stirred at -78 °C for 2 hours before the dropwise addition of chlorodimethylsilane (3.6 mL, 32.5 mmol). The reaction was allowed to warm to room temperature and stirred for 8 hours. The reaction mixture was quenched with NH₄Cl (5 mL, sat. aq.) and the product was extracted with Et₂O (15 mL × 3). The organic layer was washed with water (20 mL), brine (20 mL), dried over MgSO₄, and then concentrated under reduced pressure (50 torr). The crude product was purified by silica column chromatography with pentane to afford **1ks** (4.02 g, 16.6 mmol, 66%). ¹H NMR (400 MHz, CDCl₃) δ 7.51-7.27 (m, 7H), 7.00 (d, *J* = 8.6 Hz, 2H), 5.08 (s, 2H), 4.41 (hept, *J* = 3.7 Hz, 1H), 0.32 (d, *J* = 3.7 Hz, 6H). HRMS (FAB) *m/z*: 241.1053 ((M+H)–H₂⁺); calc. for C₁₅H₁₇OSi: 241.1049.

4-(Dimethylsilyl)aniline (23)

 H_2N

To a solution of 4-bromoaniline (3.44 g, 20.0 mmol) in DCM (30 mL) was added triethylamine (5.6 mL, 20.0 mmol) and *N*,*N*-dimethylpyridin-4-amine (DMAP, 244.3 mg, 2.0 mmol). After the mixture was stirred for 30 min, 1,2-bis(chlorodimethylsilyl)ethane

(4.78 g, 20.0 mmol) was added in one portion. The reaction was stirred at 40 °C for 3 hours. The reaction mixture was filtrated through a pad of dry Celite quickly to remove the triethylamine hydrochloride. The resulting solution was concentrated under reduced pressure (50 torr) to afford the crude product 1-(4-bromophenyl)-2,2,5,5-tetramethyl-1,2,5-azadisilolidine. (Note: this compound is moisture sensitive and decomposed on silica.)

The crude 1-(4-bromophenyl)-2,2,5,5-tetramethyl-1,2,5-azadisilolidine in THF (30 mL) in a 100 mL round-bottom flask was cooled to -78 °C. *t*-BuLi (17.6 mL, 1.7 M in pentane) was added dropwise slowly over 15 min. The resulting mixture was stirred at -78 °C for 2 hours before the dropwise addition of chlorodimethylsilane (3.3 mL, 30.0 mmol). The reaction was allowed to warm to room temperature and stirred for 8 hours. The reaction mixture was treated with Et₂O (50 mL) to allow the inorganic salts to crash out. The suspension was filtrated through a pad of Celite and basic alumina (1:1 mixture). The resulting solution was concentrated under reduced pressure (50 torr). The crude product was loaded on silica and allowed to sit for 15 min (for removal of the nitrogen

protecting group), before purification by silica column chromatography with EtOAc / hexane (1:7) to afford **23** (1.21 g, 8.00 mmol, 40%). This compound is known (62). ¹H NMR (400 MHz, CDCl₃) δ 7.38 (d, *J* = 8.4 Hz, 2H), 6.83 (d, *J* = 8.4 Hz, 2H), 4.84 (s, 2H), 4.38 (hept, *J* = 3.9 Hz, 1H), 0.30 (d, *J* = 3.7 Hz, 6H). ¹³C NMR (100 MHz, CDCl₃) δ 144.82, 135.48, 127.64, 116.07, -3.38.

Benzyl (4-(dimethylsilyl)phenyl)carbamate (23s)



To a solution of 4-(dimethylsilyl)aniline (151.2 mg, 1.0 mmol) in DCM (3 mL) was added pyridine (162 μ L, 2.0 mmol). After the mixture was stirred for 10 min, benzyl carbonochloridate (CbzCl, 170 μ L, 1.2 mmol) was added in one portion. After stirring at room temperature overnight, the reaction mixture was washed with water

(20 mL), brine (20 mL), dried over MgSO₄, and then concentrated under reduced pressure (50 torr). The crude product was purified by silica column chromatography with EtOAc / hexane (1:7) to afford **23s** (130.4 mg, 0.457 mmol, 46%). ¹H NMR (400 MHz, CDCl₃) δ 7.50 (d, J = 8.4 Hz, 2H), 7.46-7.33 (m, 5H), 6.68 (s, 1H), 5.23 (s, 2H), 4.42 (hept, J = 3.7 Hz, 1H), 0.34 (d, J = 3.7 Hz, 6H). HRMS (FAB) m/z: 286.1274 (M+H⁺); calc. for C₁₆H₂₀O₂NSi: 286.1263.

(4-Ethynylphenyl)dimethylsilane (1m)



A solution of ((4-bromophenyl)ethynyl)trimethylsilane (5.06 g, 20.0 mmol) and K_2CO_3 (5.53 g, 40.0 mmol) in MeOH (40 mL) was stirred at room temperature for 4 hours. MeOH was removed under reduced pressure, and the crude product was washed with water (30 mL) and extracted with Et₂O (40 mL). The organic layer was dried over

 $MgSO_4$, filtrated through a pad of silica, and concentrated under reduced pressure to afford the product 1-bromo-4-ethynylbenzene (3.21 g, 17.7 mmol, 89%).

In a 100 mL round-bottom flask, 1-bromo-4-ethynylbenzene (1.81 g, 10.0 mmol) in THF (15 mL) was cooled to -78 °C. *n*-BuLi (8.0 mL, 2.5 M in hexane) was added dropwise very slowly over 30 min. The resulting mixture was stirred at -78 °C for 2 hours before the dropwise addition of a solution of chlorodimethylsilane (1.1 mL, 10.0 mmol) in THF (20 mL). The reaction was allowed to warm to room temperature and stirred for 8 hours. The reaction mixture was quenched with NH₄Cl (5 mL, sat. aq.) and the product was extracted with Et₂O (15 mL × 3). The organic layer was washed with water (20 mL), brine (20 mL), dried over MgSO₄, and then concentrated under reduced pressure (200 torr). The crude product was purified by distillation under reduced pressure (1.4 torr) at 45 °C (1.12 g, 6.99 mmol, 70%). This compound is known (*63*). ¹H NMR (400 MHz, CDCl₃) δ 7.49 (d, *J* = 3.9 Hz, 4H), 4.42 (hept, *J* = 3.8 Hz, 1H), 3.10 (s, 1H), 0.34 (d, *J* = 3.8 Hz, 6H). ¹³C NMR (100 MHz, CDCl₃) δ 138.83, 134.01, 131.43, 122.91, 83.81, 77.85, -3.77.

Dimethyl(4-vinylphenyl)silane (1n)



In a 100 mL round-bottom flask, 1-bromo-4-vinylbenzene (1.83 g, 10.0 mmol) in THF (15 mL) was cooled to -78 °C. *n*-BuLi (7.5 mL, 1.6 M in hexane) was added dropwise very slowly over 30 min. The resulting mixture was stirred at -78 °C for 2 hours before the

dropwise addition of a solution of chlorodimethylsilane (1.1 mL, 10.0 mmol) in THF (20 mL). The reaction was allowed to warm to room temperature and stirred for 8 hours. The reaction mixture was quenched with NH₄Cl (5 mL, sat. aq.) and the product was extracted with Et₂O (15 mL × 3). The organic layer was washed with water (20 mL), brine (20 mL), dried over MgSO₄, and then concentrated under reduced pressure (200 torr). The crude product was purified by silica column chromatography with pentane to afford **1n** (1.29 g, 7.95 mmol, 80%). This compound is known (64). ¹H NMR (400 MHz, CDCl₃) δ 7.51 (d, *J* = 8.1 Hz, 2H), 7.41 (d, *J* = 7.9 Hz, 2H), 6.72 (dd, *J* = 17.6, 10.9 Hz, 1H), 5.79 (dd, *J* = 17.6, 0.9 Hz, 1H), 5.27 (dd, *J* = 10.9, 0.9 Hz, 1H), 4.43 (hept, *J* = 3.7 Hz, 1H), 0.35 (d, *J* = 3.8 Hz, 6H).

Cyclohexa-2,5-dien-1-yldimethylsilane (10)



In a 100 mL round-bottom flask, cyclohexa-1,4-diene (2.3 mL, 24.0 mmol) in THF (20 mL) was cooled to -78 °C. *t*-BuLi (15.5 mL, 1.7 M in pentane) and *N*,*N*,*N*,*N*-tetramethylethane-1,2-diamine (TMEDA, 3.6 mL, 24 mmol) were added simultaneously as separate solutions, dropwise over 20 min.

The resulting mixture was allowed to warm to -45 °C in 30 min and maintained at -45 °C for another 1 hour before the dropwise addition of chlorodimethylsilane (2.6 mL, 24.0 mmol). The reaction was allowed to warm to room temperature and stirred for 1.5 hours. The reaction mixture was quenched with NH₄Cl (5 mL, sat. aq.) and the product was extracted with Et₂O (15 mL × 3). The organic layer was washed with water (20 mL), brine (20 mL), dried over MgSO₄, and then concentrated under reduced pressure (300 torr). The crude product was purified by silica column chromatography with pentane to afford **10** (1.24 g, 8.97 mmol, 38%). (Note: this compound is oxygen sensitive and very volatile; storage under argon at -20° C is recommended.) This compound is known (65). ¹H NMR (400 MHz, CDCl₃) δ 5.72-5.64 (m, 2H), 5.61-5.54 (m, 2H), 3.85 (heptd, *J* = 3.5, 1.6 Hz, 1H), 2.80-2.60 (m, 2H), 2.42-2.30 (m, 1H), 0.10 (d, *J* = 3.7 Hz, 6H). ¹³C NMR (100 MHz, CDCl₃) δ 125.91, 122.09, 29.38, 26.50, -6.31.

Dimethyl(naphthalen-2-yl)silane (1p)



In a 100 mL round-bottom flask, 2-bromonaphthalene (2.07 g, 10.0 mmol) in THF (15 mL) was cooled to -78 °C. *n*-BuLi (7.5 mL, 1.6 M in hexane) was added dropwise very slowly over 30 min. The resulting mixture was stirred at -78 °C for 2 hours before the dropwise addition of a solution of chlorodimethylsilane (1.1 mL, 10.0 mmol) in THF (20 mL).

The reaction was allowed to warm to room temperature and stirred for 8 hours. The reaction mixture was quenched with NH₄Cl (5 mL, sat. aq.) and the product was extracted with Et₂O (15 mL × 3). The organic layer was washed with water (20 mL), brine (20 mL), dried over MgSO₄, and then concentrated under reduced pressure (100 torr). The crude product was purified by silica column chromatography with pentane to afford **1p** (1.69 g, 9.07 mmol, 91%). This compound is known (66). ¹H NMR (400 MHz, CDCl₃) δ 8.05 (s, 1H), 7.87-7.82 (m, 3H), 7.61 (dd, *J* = 8.1, 1.0 Hz, 1H), 7.51-7.48 (m, 2H), 4.56 (hept, *J* = 3.8 Hz, 1H), 0.43 (d, *J* = 3.8 Hz, 6H).

Benzofuran-2-yldimethylsilane (1q)

In a 100 mL round-bottom flask, benzofuran (1.18 g, 10.0 mmol) in THF (15 mL) was cooled to -78 °C. *n*-BuLi (7.5 mL, 1.6 M in hexane) was added dropwise slowly over 20 min. The resulting mixture was allowed to warm to -40 °C within 1 hour and maintained at -40 °C for another 2 hours before the dropwise addition of chlorodimethylsilane (1.1 mL, 10.0 mmol). The reaction was allowed to warm to room temperature and stirred for 8 hours. The reaction mixture was quenched with NH₄Cl (5 mL, sat. aq.) and the product was extracted with Et₂O (15 mL × 3). The organic layer was washed with water (20 mL), brine (20 mL), dried over MgSO₄, and then concentrated under reduced pressure (100 torr). The crude product was purified by silica column chromatography with Et₂O / pentane (1:30) to afford **1q** (1.08 g, 6.13 mmol, 61%). This compound is known (67). ¹H NMR (400 MHz, CDCl₃) δ 7.59 (ddd, J = 7.1, 1.3, 0.7 Hz, 1H,), 7.52 (dd, J = 8.5, 0.5 Hz, 1H), 7.30 (ddd, J = 8.5, 7.5, 1.0 Hz, 1H), 7.22 (ddd, J = 7.5, 7.5, 1.0 Hz, 1H), 7.04 (d, J = 0.7 Hz, 1H), 4.52 (hept, J = 3.8 Hz, 1H), 0.44 (d, J = 3.8 Hz, 6H).

Benzothiophen-2-yldimethylsilane (1r)

In a 100 mL round-bottom flask, benzothiophene (1.34 g, 10.0 mmol) in THF (15 mL) was cooled to -78 °C. *n*-BuLi (7.5 mL, 1.6 M in hexane) was added dropwise slowly over 20 min. The resulting mixture was allowed to warm to -40 °C within 1 hour and maintained at -40 °C for another 2 hours before the dropwise addition of chlorodimethylsilane (1.1 mL, 10.0 mmol). The reaction was allowed to warm to room temperature and stirred for 8 hours. The reaction mixture was quenched with NH₄Cl (5 mL, sat. aq.) and the product was extracted with Et₂O (15 mL × 3). The organic layer was washed with water (20 mL), brine (20 mL), dried over MgSO₄, and then concentrated under reduced pressure (100 torr). The crude product was purified by silica column chromatography with pentane to afford **1r** (1.76 g, 9.14 mmol, 91%). ¹H NMR (400 MHz, CDCl₃) δ 7.92-7.86 (m, 1H), 7.86-7.79 (m, 1H), 7.53 (s, 1H), 7.38-7.30 (m, 2H), 4.63 (hept, J = 3.7 Hz, 1H), 0.46 (d, J = 3.7 Hz, 6H). ¹³C NMR (100 MHz, CDCl₃) δ 143.87, 141.08, 138.57, 132.18, 124.47, 124.20, 123.63, 122.32, -2.93. HRMS (FAB) *m/z*: 191.0354 (M–H⁻); calc. for C₁₀H₁₂SSi: 191.0351.

V. Synthesis and Characterization of Authentic Organosilicon Products

Racemic standard references of organosilicon products were prepared using rhodium-catalyzed Si–H insertion reactions, following the general procedure described below:

General procedure for rhodium-catalyzed Si–H insertion

To a 20 mL vial or 25 mL flask was added silane (1.0 mmol, 1.0 equiv.), $Rh_2(OAc)_4$ (4.4 mg, 1 mol%) and DCM (5 mL). The mixture was cooled to -78 °C, after which diazo compound (1.0 mmol, 1.0 equiv.) was added dropwise to the solution. The reaction was allowed to slowly warm up to room temperature in 8 hours and stirred at room temperature for another 4 hours. Evaporation of the organic solvent and purification by silica column chromatography using EtOAc and hexane as eluents afforded organosilicon products **3-20** in 20-70% yields.

Organosilicon compounds **21** and **22** were prepared by rhodium-catalyzed Si–H insertion between Me-EDA and the corresponding *O*- or *N*-protected silane (**1ks** or **23s**) to give **21a** or **22a**, followed by deprotection under standard palladium-catalyzed hydrogenation condition (10% Pd/C in ethanol under H₂ atmosphere at room temperature for 16 hours). The hydrogenation reaction afforded **21** and **22** in 96% and 88% yield, respectively.

Ethyl 2-(dimethyl(phenyl)silyl)propanoate (3)



¹H NMR (400 MHz, CDCl₃) δ 7.55-7.46 (m, 2H), 7.43-7.32 (m, 3H), 4.02 (q, *J* = 7.1 Hz, 2H), 2.25 (q, *J* = 7.1 Hz, 1H), 1.17-1.11 (m, 6H), 0.37 (d, *J* = 0.6 Hz, 6H). ¹³C NMR (100 MHz, CDCl₃) δ 176.13, 136.42, 133.99, 129.57, 127.92, 59.96, 30.14, 14.43, 11.42, -3.92, -4.77. HRMS (FAB) *m/z*: 236.1234 (M⁺); calc. for C₁₃H₂₀O₂Si: 236.1233.

Ethyl 2-(dimethyl(*p*-tolyl)silyl)propanoate (4)



¹H NMR (400 MHz, CDCl₃) δ 7.40 (d, J = 7.9 Hz, 2H), 7.18 (d, J = 7.5 Hz, 2H), 4.03 (q, J = 7.1 Hz, 2H), 2.35 (s, 3H), 2.24 (q, J = 7.1 Hz, 1H), 1.19-1.11 (m, 6H), 0.35 (d, J = 0.4 Hz, 6H). ¹³C NMR (100 MHz, CDCl₃) δ 176.21, 139.51, 134.04, 132.72, 128.77, 59.94, 30.22, 21.63, 14.47, 11.44, -3.75, -4.75. ²⁹Si NMR (79 MHz, CDCl₃) δ 0.19. HRMS (FAB) m/z: 250.1391 (M⁺); calc. for C₁₄H₂₂O₂Si: 250.1389.

Ethyl 2-(dimethyl(4-methoxyphenyl)silyl)propanoate (5)



¹H NMR (400 MHz, CDCl₃) δ 7.43 (d, *J* = 8.6 Hz, 2H), 6.91 (d, *J* = 8.6 Hz, 2H), 4.03 (q, *J* = 7.1 Hz, 2H), 3.81 (s, 3H), 2.22 (q, *J* = 7.1 Hz, 1H), 1.20-1.10 (m, 6H), 0.35 (s, 6H). ¹³C NMR (100 MHz, CDCl₃) δ 176.24, 160.84, 135.49, 127.18, 113.70, 59.93, 55.18, 30.37, 14.49, 11.45, -3.72, -4.58. ²⁹Si NMR (79 MHz, CDCl₃) δ - 0.10. HRMS (FAB) *m/z*: 266.1332 (M ⁺); calc. for C₁₄H₂₂O₃Si: 266.1338.

Ethyl 2-(dimethyl(4-chlorophenyl)silyl)propanoate (6)



¹H NMR (400 MHz, CDCl₃) δ 7.43 (d, *J* = 8.4 Hz, 2H), 7.34 (d, *J* = 8.3, 1.9 Hz, 2H), 4.02 (q, *J* = 7.1 Hz, 2H), 2.23 (q, *J* = 7.1 Hz, 1H), 1.17-1.11 (m, 6H), 0.36 (s, 6H). ¹³C NMR (100 MHz, CDCl₃) δ 175.87, 135.98, 135.36, 134.75, 128.19, 60.06, 30.03, 14.45, 11.38, -4.02, -4.61. ²⁹Si NMR (79 MHz, CDCl₃) δ 0.59. HRMS (FAB) *m*/*z*: 270.0847 (M⁺); calc. for C₁₃H₁₉O₂SiCl: 270.0843.

Ethyl 2-(dimethyl(4-trifloromethylphenyl)silyl)propanoate (7)



¹H NMR (400 MHz, CDCl₃) δ 7.61 (d, J = 8.4 Hz, 4H), 4.01 (qd, J = 7.2, 0.7 Hz, 2H), 2.27 (q, J = 7.2 Hz, 1H), 1.16 (d, J = 7.2 Hz, 3H), 1.12 (t, J = 7.1 Hz, 3H), 0.40 (d, J = 1.3 Hz, 6H). ¹³C NMR (100 MHz, CDCl₃) δ 141.49, 134.32, 131.52 (q, $J_{C-F} = 32.3$ Hz), 124.26 (q, $J_{C-F} = 272.2$ Hz), 124.43 (q, $J_{C-F} = 3.8$ Hz), 60.16, 29.96, 14.39, 11.39, -4.18, -4.65. ²⁹Si NMR (79 MHz, CDCl₃) δ 1.01. HRMS (FAB) *m/z*: 304.1119 (M⁺); calc. for C₁₄H₁₉O₂SiF₃: 304.1106.

Ethyl 2-(dimethyl(4-chloromethylphenyl)silyl)propanoate (8)



¹H NMR (400 MHz, CDCl₃) δ 7.51 (d, J = 8.0 Hz, 2H), 7.38 (d, J = 7.9 Hz, 2H), 4.58 (s, 2H), 4.02 (q, J = 7.1 Hz, 2H), 2.25 (q, J = 7.1 Hz, 1H), 1.18-1.08 (m, 6H), 0.37 (d, J = 1.2 Hz, 6H). ¹³C NMR (100 MHz, CDCl₃) δ 175.98, 138.72, 136.92, 134.44, 128.02, 60.02, 46.23, 30.05, 14.44, 11.41, -3.97, -4.70. ²⁹Si NMR (79 MHz, CDCl₃) δ 0.59. HRMS (FAB) *m*/*z*: 283.0927 ((M+H)–H₂ ⁺); calc. for C₁₄H₂₀O₂SiCl: 283.0921.

Methyl 4-((1-ethoxy-1-oxopropan-2-yl)dimethylsilyl)benzoate (9)



¹H NMR (400 MHz, CDCl₃) δ 8.00 (d, *J* = 8.3 Hz, 2H), 7.58 (d, *J* = 8.3 Hz, 2H), 4.01 (qd, *J* = 7.1, 0.9 Hz, 2H), 3.92 (s, 3H), 2.27 (q, *J* = 7.2 Hz, 1H), 1.19-1.05 (m, 6H), 0.39 (d, *J* = 1.4 Hz, 6H). ¹³C NMR (100 MHz, CDCl₃) δ 175.77, 167.27, 142.78, 134.01, 131.00, 128.64, 60.07, 52.31, 29.88, 14.44, 11.37, -4.13, -4.71. ²⁹Si NMR (79 MHz, CDCl₃) δ 0.98. HRMS (FAB) *m/z*: 294.1276 (M⁺); calc. for C₁₅H₂₂O₄Si: 294.1287.

Ethyl 2-((4-(dimethylcarbamoyl)phenyl)dimethylsilyl)propanoate (10)



¹H NMR (400 MHz, CDCl₃) δ 7.54 (d, *J* = 8.0 Hz, 2H), 7.39 (d, *J* = 8.0 Hz, 2H), 4.03 (q, *J* = 7.1 Hz, 2H), 3.11 (s, 3H), 2.97 (s, 3H), 2.26 (q, *J* = 7.2 Hz, 1H), 1.20-1.10 (m, 6H), 0.37 (s, 6H). ¹³C NMR (100 MHz, CDCl₃) δ 175.94, 171.59, 138.39, 137.38, 134.01, 126.37, 60.05, 39.67, 35.45, 29.95, 14.49, 11.41, -3.92, -4.75. HRMS (FAB) *m/z*: 308.1689 (M+H⁺); calc. for C₁₆H₂₆NO₃Si: 308.1682.

Ethyl 2-((naphthalen-2-yl)dimethylsilyl)propanoate (11)



¹H NMR (400 MHz, CDCl₃) δ 8.00 (s, 1H), 7.87-7.80 (m, 3H), 7.57 (dd, J = 8.2, 0.8 Hz, 1H), 7.53-7.46 (m, 2H), 4.02 (q, J = 7.1 Hz, 2H), 2.34 (q, J = 7.1 Hz, 1H), 1.18 (d, J = 7.1 Hz, 3H), 1.11 (t, J = 7.1 Hz, 3H), 0.46 (d, J = 2.0 Hz, 6H). ¹³C NMR (100 MHz, CDCl₃) δ 176.14, 134.88, 133.99, 133.92, 132.92, 130.03, 128.23, 127.84, 127.16, 126.70, 126.15, 60.01, 30.18, 14.44, 11.50, -3.76, -4.71. ²⁹Si NMR (79 MHz, CDCl₃) δ 0.76. HRMS (FAB) m/z: 286.1395 (M⁺); calc. for C₁₇H₂₂O₂Si: 286.1389.

Ethyl 2-((benzofuran-2-yl)dimethylsilyl)propanoate (12)



¹H NMR (400 MHz, CDCl₃) δ 7.58 (ddd, J = 7.7, 1.3, 0.7 Hz, 1H), 7.50 (dd, J = 8.2, 0.9 Hz, 1H), 7.29 (ddd, J = 8.4, 7.2, 1.4 Hz, 1H), 7.21 (ddd, J = 8.2, 7.8, 1.0 Hz, 1H), 7.05 (d, J = 1.0 Hz, 1H), 4.07 (qd, J = 7.2, 1.2 Hz, 2H), 2.38 (q, J = 7.2 Hz, 1H), 1.24 (d, J = 7.1 Hz, 3H), 1.14 (t, J = 7.1 Hz, 3H), 0.44 (d, J = 1.9 Hz, 6H). ¹³C NMR (100 MHz, CDCl₃) δ 175.57, 159.85, 158.31, 127.82, 124.84, 122.58, 121.29, 117.93, 111.48, 60.22, 29.42, 14.42, 11.24, -4.32, -5.05. ²⁹Si NMR (79)

MHz, CDCl₃) δ –5.09. HRMS (FAB) *m*/*z*: 276.1182 (M⁺); calc. for C₁₅H₂₀O₃Si: 276.1169.

Ethyl 2-(benzothiophen-2-yldimethylsilyl)propanoate (13)



Ethyl 2-(benzyldimethylsilyl)propanoate (14)



¹H NMR (400 MHz, CDCl₃) δ 7.25-7.17 (m, 2H), 7.11-7.06 (m, 1H), 7.06-6.96 (m, 2H), 4.24-4.02 (m, 2H), 2.24-2.14 (m, 2H), 2.10 (q, *J* = 7.1 Hz, 1H), 1.26 (t, *J* = 7.1 Hz, 3H), 1.19 (d, *J* = 7.1 Hz, 3H), 0.06- -0.12 (m, *J* = 0.5 Hz, 6H). ¹³C NMR (100 MHz, CDCl₃) δ 176.19, 139.25, 128.44, 128.42, 124.44, 60.04, 28.95, 24.03, 14.66, 11.17, -4.76, -4.82. ²⁹Si NMR (79 MHz, CDCl₃) δ 5.96. HRMS (FAB) *m*/*z*: 251.1459 (M+H⁺); calc. for C₁₄H₂₃O₂Si: 251.1467.

Ethyl 2-(cyclohexa-2,5-dien-1-yldimethylsilyl)propanoate (15)



¹H NMR (400 MHz, CDCl₃) δ 5.81-5.52 (m, 4H), 4.17-4.05 (m, 2H), 2.81-2.56 (m, 2H), 2.48-2.36 (m, 1H), 2.19 (q, *J* = 7.1 Hz, 1H), 1.24 (t, *J* = 7.1 Hz, 3H), 1.21 (d, *J* = 7.2 Hz, 3H), 0.08 (d, *J* = 4.4 Hz, 6H). ¹³C NMR (100 MHz, CDCl₃) δ 176.24, 125.63, 125.53, 122.65, 122.52, 60.05, 29.47, 28.28, 26.52, 14.59, 11.33, -6.35, -6.70. HRMS (FAB) *m/z*: 237.1313 ((M+H)-H₂⁺); calc. for C₁₃H₂₁O₂Si: 237.1311.

Ethyl 2-((4-ethynylphenyl)dimethylsilyl)butanoate (16)



¹H NMR (400 MHz, CDCl₃) δ 7.53-7.42 (m, 4H), 4.13-3.93 (m, 2H), 3.11 (s, 1H), 2.08 (dd, J = 11.7, 3.1 Hz, 1H), 1.78 (ddq, J = 14.2, 11.7, 7.1 Hz, 1H), 1.40 (dqd, J = 13.8, 7.3, 3.1 Hz, 1H), 1.13 (t, J = 7.1 Hz, 3H), 0.89 (t, J = 7.2 Hz, 3H), 0.35 (d, J = 4.1 Hz, 6H). ¹³C NMR (100 MHz, CDCl₃) δ 175.05, 137.86, 133.87, 131.36, 123.17, 83.71, 78.06, 59.95, 39.53, 20.60, 15.13, 14.47, – 3.96, –4.56. HRMS (FAB) *m/z*: 273.1316 ((M+H)–H₂⁺); calc. for C₁₆H₂₁O₂Si: 273.1311.

Isopropyl 2-((4-ethynylphenyl)dimethylsilyl)propanoate (17)



¹H NMR (400 MHz, CDCl₃) δ 7.47 (app s, 4H), 4.91 (hept, J = 6.3 Hz, 1H), 3.11 (s, 1H), 2.21 (q, J = 7.1 Hz, 1H), 1.16 (d, J = 6.3 Hz, 3H), 1.13 (d, J = 7.2 Hz, 3H), 1.08 (d, J = 6.3 Hz, 3H), 0.36 (d, J = 3.0 Hz, 6H). ¹³C NMR (100 MHz, CDCl₃) δ 175.39, 137.82, 133.93, 131.36, 123.15, 83.73, 78.03, 67.33, 30.02, 22.16, 22.01, 11.45, -3.95, -4.69. HRMS (FAB) *m/z*: 275.1479 (M+H⁺); calc. for C₁₆H₂₃O₂Si: 275.1467.

Ethyl 2-((3,4-dihydro-2H-pyran-6-yl)dimethylsilyl)propanoate (18)



¹H NMR (400 MHz, CDCl₃) δ 5.02 (t, *J* = 3.8, 1H), 4.16-4.02 (m, 2H), 3.90 (dd, *J*=5.8, 4.9 Hz, 2H), 2.17 (q, *J* = 7.1 Hz, 1H), 2.05-1.98 (m, 2H), 1.87-1.78 (m, 2H), 1.23 (t, *J* = 7.1 Hz, 3H), 1.18 (d, *J* = 7.1 Hz, 3H), 0.14 (d, *J* = 8.6 Hz, 6H). ¹³C NMR (100 MHz, CDCl₃) δ 176.26, 157.26, 112.83, 65.77, 59.91, 28.78, 22.79, 20.90, 14.61, 11.24, -4.96, -5.98. ²⁹Si NMR (79 MHz, CDCl₃) δ -4.02. HRMS (FAB) *m*/*z*: 243.1410 (M+H⁺); calc. for C₁₂H₂₃O₃Si: 243.1417.

Ethyl 2-((4-vinylphenyl)dimethylsilyl)propanoate (19)



¹H NMR (400 MHz, CDCl₃) δ 7.47 (d, *J* = 8.1 Hz, 2H), 7.40 (d, *J* = 8.0 Hz, 2H), 6.71 (dd, *J* = 17.6, 10.9 Hz, 1H), 5.79 (dd, *J* = 17.6, 0.9 Hz, 1H), 5.27 (dd, *J* = 10.9, 1.0 Hz, 1H), 4.03 (q, *J* = 7.1 Hz, 2H), 2.25 (q, *J* = 7.1 Hz, 1H), 1.16-1.13 (m, 6H), 0.37 (s, 6H). ¹³C NMR (100 MHz, CDCl₃) δ 176.09, 138.68, 136.85, 136.00, 134.26, 125.69, 114.65, 59.99, 30.14, 14.46, 11.42, -3.87, -4.75. ²⁹Si NMR (79 MHz,

CDCl₃) δ 0.26. HRMS (FAB) *m*/*z*: 262.1384 (M⁺); calc. for C₁₅H₂₂O₂Si: 262.1389.

Ethyl 2-((4-ethynylphenyl)dimethylsilyl)propanoate (20)



¹H NMR (400 MHz, CDCl₃) δ 7.50-7.44 (m, 4H), 4.01 (qd, J = 7.2, 0.6 Hz, 2H), 3.11 (s, 1H), 2.24 (q, J = 7.1 Hz, 1H), 1.17-1.10 (m, 6H), 0.37 (d, J = 0.7 Hz, 6H). ¹³C NMR (100 MHz, CDCl₃) δ 175.89, 137.69, 133.87, 131.36, 123.20, 83.70, 78.08, 60.04, 29.98, 14.44, 11.38, -4.10, -4.75. ²⁹Si NMR (79 MHz, CDCl₃) δ 0.79. HRMS (FAB) m/z: 259.1153 ((M+H)–H₂⁺); calc. for C₁₅H₁₉O₂Si: 259.1154.

Ethyl 2-((4-(benzyloxy)phenyl)dimethylsilyl)propanoate (21a)



¹H NMR (400 MHz, CDCl₃) δ 7.46-7.30 (m, 7H), 7.05-6.93 (m, 2H), 5.08 (s, 2H), 4.02 (q, *J* = 7.1 Hz, 2H), 2.22 (q, *J* = 7.1 Hz, 1H), 1.20-1.10 (m, 6H), 0.35 (s, 6H). ¹³C NMR (100 MHz, CDCl₃) δ 176.22, 160.07, 137.01, 135.51, 128.74, 128.13, 127.61, 127.52, 114.56, 69.87, 59.93, 30.35, 14.48, 11.45, -3.75, -4.57. HRMS (FAB) *m/z*: 342.1665 (M⁺); calc. for C₂₀H₂₆O₃Si: 342.1651.

Ethyl 2-((4-hydroxyphenyl)dimethylsilyl)propanoate (21)



¹H NMR (400 MHz, CDCl₃) δ 7.36 (d, *J* = 8.5 Hz, 2H), 6.81 (d, *J* = 8.6 Hz, 2H), 5.49 (s, 1H), 4.03 (qd, *J* = 7.1, 0.6 Hz, 2H), 2.23 (q, *J* = 7.1 Hz, 1H), 1.19-1.12 (m, 6H), 0.35 (d, *J* = 1.2 Hz, 6H). ¹³C NMR (100 MHz, CDCl₃) δ 176.67, 157.18, 135.68, 126.90, 115.16, 60.12, 30.51, 14.46, 11.43, -3.83, -4.33. HRMS (FAB) *m*/*z*: 252.1175 (M⁺); calc. for C₁₃H₂₀O₃Si: 252.1182.

Ethyl 2-((4-(((benzyloxy)carbonyl)amino)phenyl)dimethylsilyl)propanoate (22a)



¹H NMR (400 MHz, CDCl₃) δ 7.49-7.30 (m, 9H), 6.70 (s, 1H), 5.20 (s, 2H), 4.02 (q, *J* = 7.1 Hz, 2H), 2.22 (q, *J* = 7.1 Hz, 1H), 1.18-1.10 (m, 6H), 0.35 (s, 6H). ¹³C NMR (100 MHz, CDCl₃) δ 176.13, 153.24, 139.07, 136.06, 134.98, 130.78, 128.78, 128.56, 128.48, 117.96, 67.24, 59.98, 30.21, 14.49, 11.41, -3.83, -4.68. HRMS (FAB) *m*/*z*: 385.1710 (M⁺); calc. for C₂₁H₂₇NO₄Si: 385.1709.

Ethyl 2-((4-aminophenyl)dimethylsilyl)propanoate (22)



¹H NMR (400 MHz, CDCl₃) δ 7.29 (d, *J* = 8.4 Hz, 2H), 6.70 (d, *J* = 8.4 Hz, 2H), 4.03 (q, *J* = 7.1 Hz, 2H), 3.64 (br s, 2H), 2.20 (q, *J* = 7.1 Hz, 1H), 1.16 (t, *J* = 7.1 Hz, 3H), 1.13 (d, *J* = 7.1 Hz, 3H), 0.32 (s, 6H). ¹³C NMR (100 MHz, CDCl₃) δ 176.40, 147.40, 135.32, 124.37, 114.82, 59.87, 30.47, 14.48, 11.44, -3.68, -4.68. HRMS (FAB) *m/z*: 251.1339 (M⁺); calc. for C₁₃H₂₁NO₂Si: 251.1342. [α_D]²⁵ = +46.1 (*c* 0.505 in cyclohexane, 98% *ee*).

VI. Preparative-Scale Whole-Cell Biocatalytic Reaction



 $HB_{amp/chlor}$ (50 mL) in a 250 mL flask was inoculated with an overnight culture (1 mL, LB_{amp/chlor}) of recombinant E. cloni[®] EXPRESS BL21(DE3) cells containing a pET22(b)+ plasmid encoding Rma cyt c V75T M100D M103D, and the pEC86 plasmid. The culture was shaken at 37 °C and 250 rpm (no humidity control) until the OD₆₀₀ was 0.6 (approximately 2 hours). The culture was placed on ice for 30 minutes, and IPTG and 5-ALA were added to final concentrations of 20 µM and 200 µM respectively. The incubator temperature was reduced to 20 °C, and the culture was allowed to shake for 20 hours at 140 rpm. Cells were pelleted by centrifugation (4 °C, 5 min, 4,000xg), resuspended in M9-N buffer and adjusted to $OD_{600} = 15$. Under anaerobic conditions, to a 5 mL reaction vial was added 960 μ L whole-cell solution, 40 μ L glucose solution (250 mM in M9-N buffer), 15.8 μL 4-(dimethylsilyl)aniline (23, 0.1 mmol) and 4.9 μL Me-EDA (0.04 mmol) at room temperature. The reaction was shaken at 480 rpm. At 4 h intervals, two additional batches of whole cell (960 μ L), glucose (40 μ L) and Me-EDA $(4.9 \ \mu\text{L}, 0.04 \ \text{mmol})$ were added to the reaction. After the reaction was shaken for a total of 20 hours, the reaction mixture was divided between four 2 mL microcentrifuge tubes, and 0.6 mL EtOAc was added to each tube. The reaction mixtures were subjected to vortex (30 seconds) and centrifugation (14,000 rpm, 7 min) to completely separate the organic and aqueous layers. After removal of the organic layers, two addition rounds of extraction were carried out. The combined organic layers were dried over anhydrous Na₂SO₄, concentrated, and purified by silica column chromatography with EtOAc / hexane (1:4 to 1:2) to afford pure organosilicon product 22 (17.6 mg, 0.0700 mmol, 70% yield), together with recovered silane 23 (4.0 mg, 0.0265 mmol, 26.5%). The stereoselectivity of the product was determined as 98% ee by chiral SFC. The protein concentration of $OD_{600} = 15$ whole-cell solution was determined to be 7.43 μ M by the ferrous assay after cell lysis by sonication. The total turnover number for this reaction was 3410.

VII. GC Standard Curves for Organosilicon Products

The analysis of product formation in enzymatic reactions was performed based on GC standard curves. The general procedure for preparing analytical samples for GC standard curves is described below:

Sample preparation for GC standard curves

Stock solutions of chemically synthesized organosilicon products at various concentrations (20 to 200 mM in MeCN) were prepared. To a microcentrifuge tube were added 340 μ L M9-N buffer, 40 μ L Na₂S₂O₄ (100 mM in dH₂O), 20 μ L organosilicon product, 20 μ L internal standard (20 mM 2-phenylethanol in cyclohexane) and 1 mL cyclohexane. The mixture was vortexed (10 seconds, 3 times) then centrifuged (14,000xg, 5 min) to completely separate the organic and aqueous layers. The organic layer (750 μ L) was removed for GC analysis. All data points represent the average of duplicate runs. The standard curves plot product concentration in mM (y-axis) against the ratio of product area to internal standard area on the GC (x-axis).






































VIII. Chiral SFC Traces for Racemic and Enzymatically Synthesized Organosilicon Products

All the *ee* values of enzymatically synthesized organosilicon products were determined using chiral SFC. The chiral SFC traces for racemic and enzymatic products are shown below.

The absolute configuration of enzymatically synthesized organosilicon product **3** was determined to be (*R*) by comparing the HPLC retention times of *rac*-**3** and **3** with that reported in the literature (12). The absolute configurations of other organosilicon products were inferred by analogy, assuming the facial selectivity of the diazo reagents remains the same in the biosyntheses of compound **3**-**22**.



rac or R-3



Chiralpak IC, 3% i-PrOH in CO₂, 3.5 mL/min, 210 nm

Area% report for <i>rac</i> and <i>R</i> -3:								
rac-3			<i>R</i> -3					
Retention	Area	Aroa Va	Retention	Area	Aroo 0/2			
Time (min)	(mAU*s)	Alea 70	Time (min)	(mAU*s)	Alta 70			
2.224	842.723	50.03	2.172	753.160	100.00			
2.390	841.657	49.97	-	-				
Total	1684.380	100.00	Total	753.160	100.00			

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Area% re	port for	rac	and R -4	:
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rac-4			<i>R</i> -4		
Retention	Area	A ==== 07	Retention	Area	A mag 07
Time (min)	(mAU*s)	Alea %	Time (min)	(mAU*s)	Alea %
2.097	124486	45.44	2.142	16934.0	100.00
2.390	149493	54.56	-	-	
Total	273979	100.00	Total	16934.0	100.00





Area% report for *rac* and *R*-**5**:

inea /o report r					
rac-5			<i>R</i> -5		
Retention	Area	Area 07-	Retention	Area	Aroo 07-
Time (min)	(mAU*s)	Alea 70	Time (min)	(mAU*s)	Alea %
3.869	8651.07	49.14	3.873	1422.48	100.00
4.349	8955.03	50.86	-	-	
Total	17606.10	100.00	Total	1422.48	100.00





Area%	report	for	rac	and R	- 6 :
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rac-6			<i>R</i> -6		
Retention	Area	Aroo 02	Retention	Area	Aroo 07-
Time (min)	(mAU*s)	Alta 70	Time (min)	(mAU*s)	Alta 70
2.510	4898.62	49.48	2.477	455.973	100.00
2.746	5001.36	50.52	-	-	
Total	9899.98	100.00	Total	455.973	100.00



Chiralcel OD-H, 2% *i*-PrOH in CO₂, 2.5 mL/min, 210 nm



Area% rep	port for <i>rac</i>	and <i>R</i>-7 :
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rac-7			<i>R</i> -7		
Retention	Area	Area 02	Retention	Area	Aroo 07-
Time (min)	(mAU*s)	Alta 70	Time (min)	(mAU*s)	Alta 70
2.186	3895.65	49.54	2.182	1738.84	99.61
2.431	3967.58	50.46	2.402	6.80	0.39
Total	7863.23	100.00	Total	1745.65	100.00





Area% report for *rac* and *R*-8:

rac-8			<i>R</i> -8		
Retention	Area	A #20 07-	Retention	Area	Area 07-
Time (min)	(mAU*s)	Alea %	Time (min)	(mAU*s)	Alea 70
3.406	1698.74	49.77	3.409	2377.97	99.58
3.914	1714.41	50.23	3.912	10.15	0.42
Total	3413.15	100.00	Total	2388.12	100.00



Chiralcel OD-H, 5% *i*-PrOH in CO₂, 2.5 mL/min, 210 nm



Area% report for	or <i>rac</i> and R-9 :				
rac-9			<i>R</i> -9		
Retention	Area	Aroa Va	Retention	Area	Aroo %
Time (min)	(mAU*s)	Alea 70	Time (min)	(mAU*s)	Alta 70
3.517	1773.21	50.00	3.507	582.064	100.00
3.824	1772.88	50.00	-	-	
Total	3546.09	100.00	Total	582.064	100.00

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Chiralpak IC, 30% *i*-PrOH in CO_2 , 2.5 mL/min, 210 nm



Area%	report	for	rac	and	R -	1():
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<i>rac</i> -10			<i>R</i> -10		
Retention	Area	Aroo 07-	Retention	Area	Area 07-
Time (min)	(mAU*s)	Alea %	Time (min)	(mAU*s)	Alea 70
6.693	2779.93	49.38	6.665	987.666	99.63
7.000	2849.58	50.62	7.005	3.594	0.36
Total	5629.50	100.00	Total	991.260	100.00



Chiralpak IC, 7% *i*-PrOH in CO₂, 3.5 mL/min, 210 nm



Area% report for *rac* and *R*-11:

<i>rac</i> -11			<i>R</i> -11		
Retention	Area	$\Lambda mag 07$	Retention	Area	A ==== 07
Time (min)	(mAU*s)	Alea %	Time (min)	(mAU*s)	Alea %
2.735	8677.04	47.45	2.760	3675.15	97.51
3.183	9609.04	52.55	3.215	93.95	2.49
Total	18286.1	100.00	Total	3769.10	100.00





Area% report for *rac* and *R*-12:

<i>rac</i> -12			<i>R</i> -12		
Retention	Area	Area 07-	Retention	Area	Aroo 07-
Time (min)	(mAU*s)	Alea 70	Time (min)	(mAU*s)	Alea %
3.101	7145.58	49.41	3.034	1490.62	99.16
3.624	7315.99	50.59	3.549	12.69	0.84
Total	14461.6	100.00	Total	1503.31	100.00





rac-13			<i>R</i> -13		
Retention	Area	Aroa Va	Retention	Area	Aroa Va
Time (min)	(mAU*s)	Alea 70	Time (min)	(mAU*s)	Alta 70
2.751	7136.78	48.49	2.764	801.009	98.78
3.209	7581.08	51.51	3.219	9.889	1.22
Total	14717.9	100.00	Total	810.898	100.00



Chiralpak AD-H, 3% *i*-PrOH in CO₂, 2.5 mL/min, 210 nm



Area% report for *rac* and *R*-14:

<i>rac</i> -14			<i>R</i> -14		
Retention	Area	Area 07-	Retention	Area	Area 07-
Time (min)	(mAU*s)	Alea 70	Time (min)	(mAU*s)	Alea %
2.142	1384.73	49.85	2.191	1991.26	99.62
2.351	1393.07	50.15	2.402	7.58	0.38
Total	2777.80	100.00	Total	1998.84	100.00





rac-15			<i>R</i> -15		
Retention	Area	Aroa Va	Retention	Area	Aron 0/2
Time (min)	(mAU*s)	Alea 70	Time (min)	(mAU*s)	Alea 70
2.356	361.685	50.30	2.426	390.115	99.45
2.500	357.384	49.70	2.556	2.125	0.54
Total	719.070	100.00	Total	392.240	100.00





Area% repo	rt for <i>rac</i>	and <i>R</i>-16 :
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<i>rac</i> -16			<i>R</i> -16		
Retention	Area	A ==== 07	Retention	Area	A ==== 07
Time (min)	(mAU*s)	Alea %	Time (min)	(mAU*s)	Alea %
2.602	4276.04	49.44	2.607	2426.65	100.00
3.526	4373.23	50.56			
Total	8649.28	100.00	Total	2426.65	100.00



Chiralpak IC, 3% *i*-PrOH in CO₂, 2.5 mL/min, 210 nm



Area% report for *rac* and *R*-17:

<i>rac</i> -17			<i>R</i> -17		
Retention	Area	Aroo 0%	Retention	Area	Aroo Va
Time (min)	(mAU*s)	Alta 70	Time (min)	(mAU*s)	Alea 70
3.808	1442.74	49.78	3.829	2089.15	100.00
4.074	1455.49	50.22			
Total	2898.23	100.00	Total	2089.15	100.00





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<i>rac</i> -18			<i>R</i> -18		
Retention	Area	A #22 07	Retention	Area	$\Lambda mag 07$
Time (min)	(mAU*s)	Alea %	Time (min)	(mAU*s)	Area %
2.680	1366.02	50.54	2.628	813.657	99.63
2.887	1336.57	49.46	2.922	2.984	0.37
Total	2702.59	100.00	Total	816.641	100.00





Area% report for *rac* and *R*-19:

rac-19			<i>R</i> -19		
Retention	Area	Aroo 07-	Retention	Area	Aroo 07-
Time (min)	(mAU*s)	Alea %	Time (min)	(mAU*s)	Alea %
2.791	6332.09	49.69	2.758	1559.90	99.13
3.001	6411.17	50.31	2.972	13.65	0.87
Total	12743.3	100.00	Total	1573.56	100.00



Chiralcel OD-H, 3% *i*-PrOH in CO₂, 2.5 mL/min, 210 nm



Area%	report	for	rac	and	R -2	20 :
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<i>rac</i> -20			<i>R</i> -20		
Retention	Area	Aroo 07-	Retention	Area	Aroo 07-
Time (min)	(mAU*s)	Alea %	Time (min)	(mAU*s)	Alea %
3.410	13436.2	48.40	3.406	1894.53	99.84
3.774	14326.4	51.60	3.715	2.97	0.16
Total	27762.6	100.00	Total	1897.50	100.00





Area%	report	for	R-21 :	
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<i>rac</i> -21			<i>R</i> -21		
Retention	Area	Aroo 02	Retention	Area	Aroo 02
Time (min)	(mAU*s)	Alea %	Time (min)	(mAU*s)	Alea %
3.853	664.781	50.19	3.883	3332.68	99.58
4.264	659.660	49.81	4.331	13.94	0.42
Total	1324.441	100.00	Total	3346.62	100.00



Chiralpak AD-H, 10% i-PrOH in CO₂, 2.5 mL/min, 210 nm



Area% report for *rac* and *R*-22:

rac-22			<i>R</i> -22		
Retention	Area	A 1000 07	Retention	Area	$\Lambda mag 07$
Time (min)	(mAU*s)	Alea %	Time (min)	(mAU*s)	Alea %
5.162	9214.41	48.50	-	-	
5.617	9783.58	51.50	5.638	195.47	100.00
Total	18998.0	100.00	Total	195.47	100.00

Preparative-scale reaction (0.1 mmol):



Area% report for *R***-22**:

<i>R</i>-22 (OD 15-whole cell)					
Retention Time (min)	Area (mAU*s)	Area %			
5.537	45.97	1.15			
5.990	3935.62	98.85			
Total	3981.59	100.00			

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X. NMR Spectra
























fl (ppm) -10





















































²⁹Si NMR (79 Hz, CDCl₃; referenced at δ –78.9 ppm using tetramethoxysilane as internal standard. This internal standard contained a small amount of impurity, which appeared at δ –86.3 ppm.)














