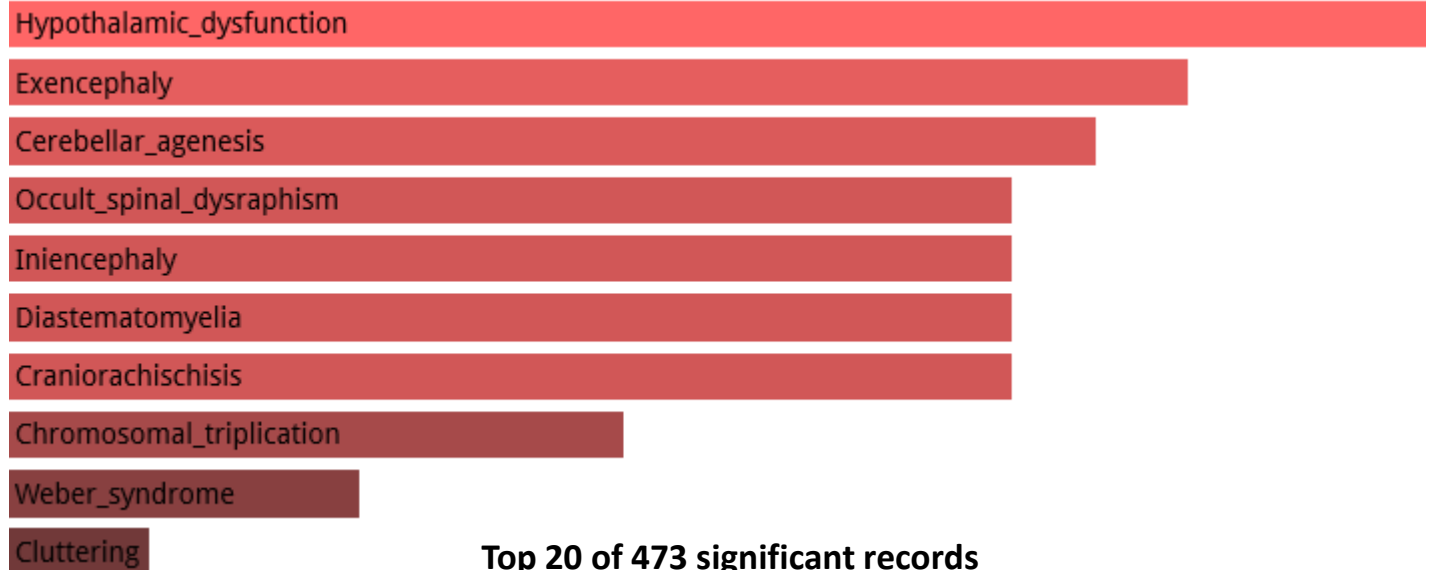


**Supplemental Figure S5.** Identification of genes

implicated in more than 1,000 records classified as human rare diseases using the Enrichr bioinformatics platform

(see Methods for details).

# Rare Diseases GeneRIF Gene Lists: 8,405 genes



Top 20 of 473 significant records

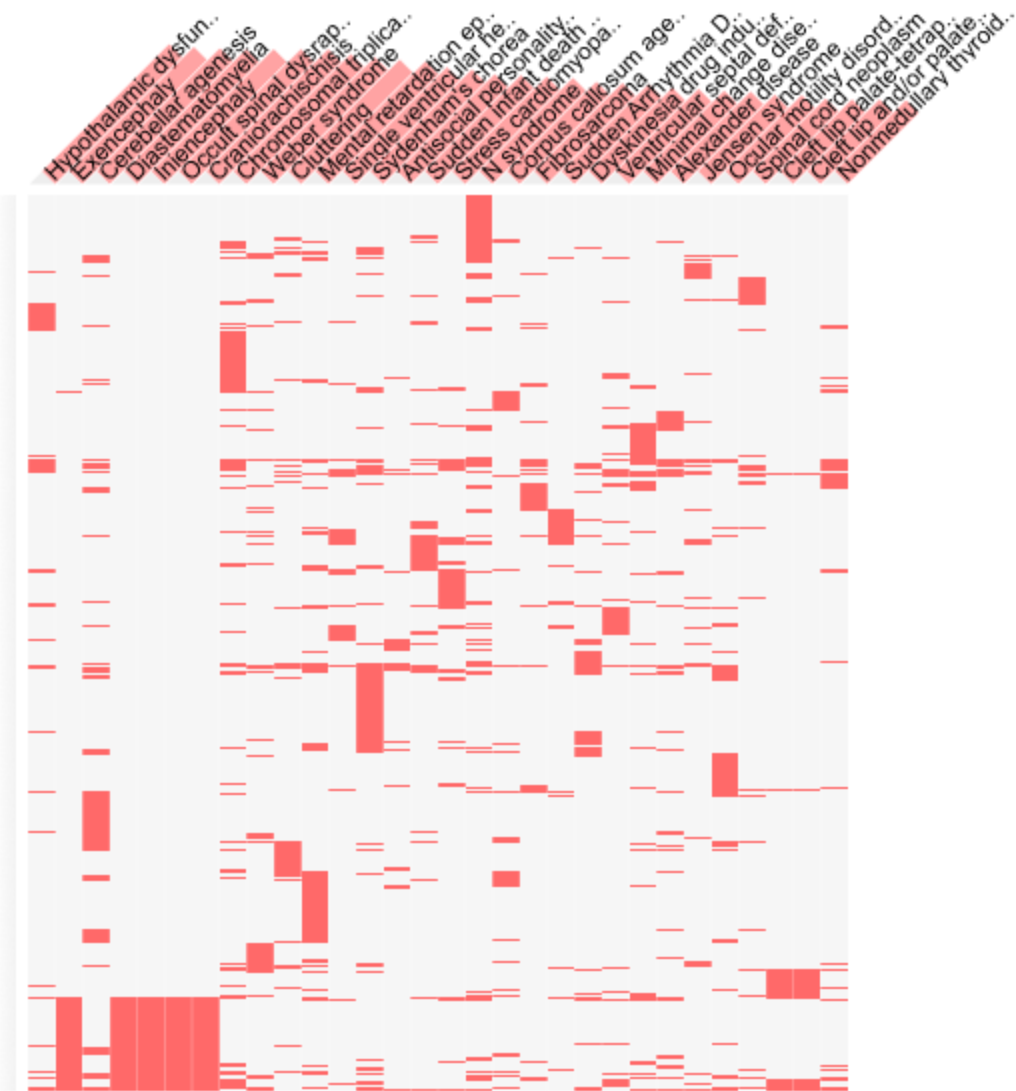
Term	Overlap	P-value	Adjusted P-value
Hypothalamic_dysfunction	90/128	8.12E-11	6.55E-08
Exencephaly	158/256	1.38E-10	6.55E-08
Cerebellar_agenesis	198/335	1.7E-10	6.55E-08
Occult_spinal_dysraphism	157/255	2.05E-10	6.55E-08
Diastematomyelia	157/255	2.05E-10	6.55E-08
Craniorachischisis	157/255	2.05E-10	6.55E-08
Iniencephaly	157/255	2.05E-10	6.55E-08
Chromosomal_triplication	201/344	4.9E-10	1.37E-07
Weber_syndrome	94/139	8.85E-10	2.2E-07
Cluttering	106/162	1.41E-09	3.16E-07
Mental_retardation_epilepsy	190/326	1.92E-09	3.9E-07
Single_ventricular_heart	73/104	5.49E-09	1.02E-06
Sydenham's_chorea	209/369	8.42E-09	1.34E-06
Chorea_minor	209/369	8.42E-09	1.34E-06
Antisocial_personality_disorder	46/59	1.93E-08	2.87E-06
Sudden_infant_death_syndrome	103/162	2.37E-08	3.31E-06
Stress_cardiomyopathy	110/176	3.16E-08	4.16E-06
N_syndrome	222/401	3.89E-08	4.83E-06
Basilar_migraine	127/210	4.85E-08	5.7E-06
Corpus_callosum_agenesis	89/138	7.92E-08	8.85E-06

# Rare Diseases GeneRIF Gene Lists: 8,405 genes

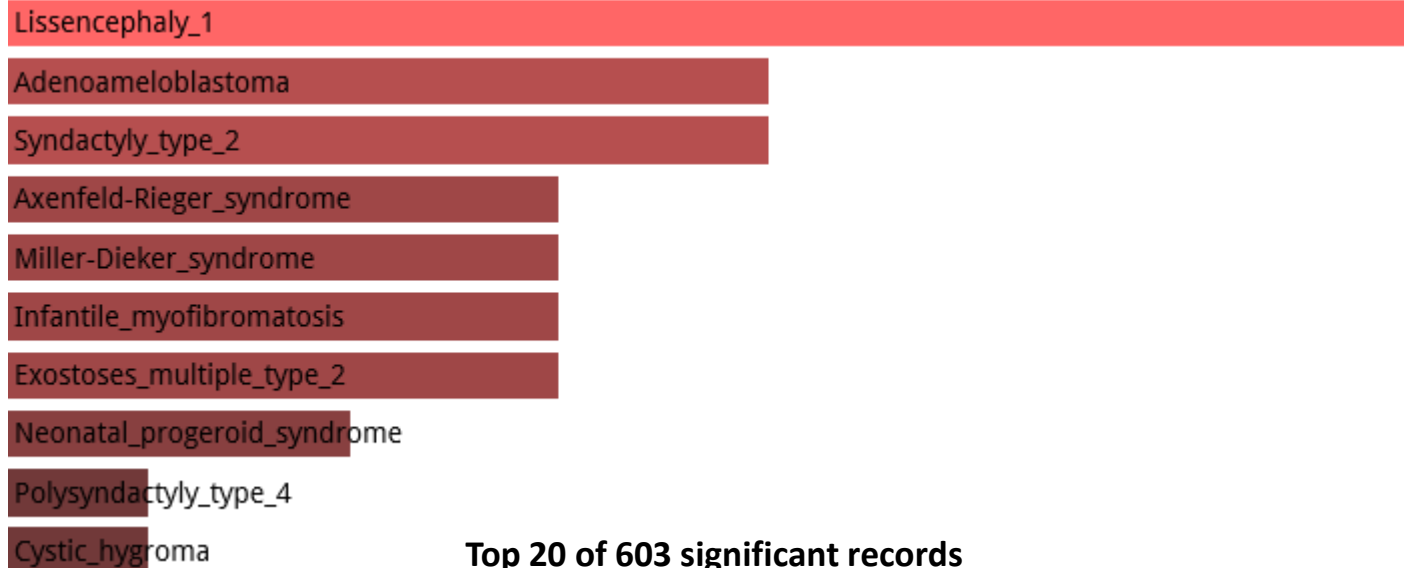


## Enriched Terms

Input Genes



# Rare Diseases GeneRIF ARCHS4 Predictions: 8,405 genes



Top 20 of 603 significant records

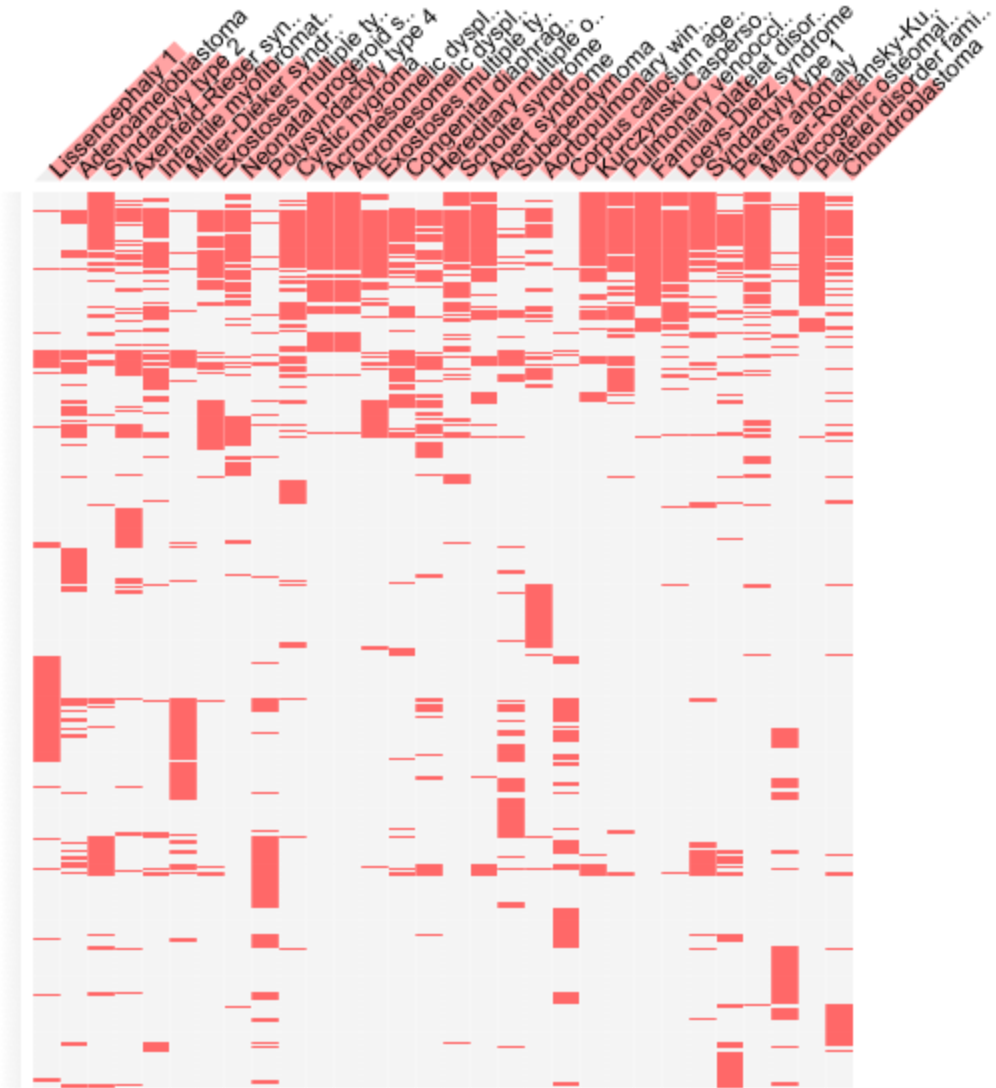
Term	Overlap	P-value	Adjusted P-value
Lissencephaly_1	154/200	4.79E-24	1.08E-20
Adenoameloblastoma	151/200	4.35E-22	3.26E-19
Syndactyly_type_2	151/200	4.35E-22	3.26E-19
Axenfeld-Rieger_syndrome	150/200	1.85E-21	5.94E-19
Miller-Dieker_syndrome	150/200	1.85E-21	5.94E-19
Exostoses_multiple_type_2	150/200	1.85E-21	5.94E-19
Infantile_myofibromatosis	150/200	1.85E-21	5.94E-19
Neonatal_progeroid_syndrome	149/200	7.69E-21	2.16E-18
Polysyndactyly_type_4	148/200	3.11E-20	6.97E-18
Cystic_hygrroma	148/200	3.11E-20	6.97E-18
Acromesomelic_dysplasia_Hunter_Thompson_type	147/200	1.22E-19	2.11E-17
Acromesomelic_dysplasia	147/200	1.22E-19	2.11E-17
Exostoses_multiple_type_1	147/200	1.22E-19	2.11E-17
Scholte_syndrome	146/200	4.69E-19	6.58E-17
Congenital_diaphragmatic_hernia	146/200	4.69E-19	6.58E-17
Hereditary_multiple_osteochondromas	146/200	4.69E-19	6.58E-17
Aortopulmonary_window	145/200	1.76E-18	1.88E-16
Corpus_callosum_agenesis	145/200	1.76E-18	1.88E-16
Apert_syndrome	145/200	1.76E-18	1.88E-16
Subependymoma	145/200	1.76E-18	1.88E-16

# Rare Diseases GeneRIF ARCHS4 Predictions: 8,405 genes

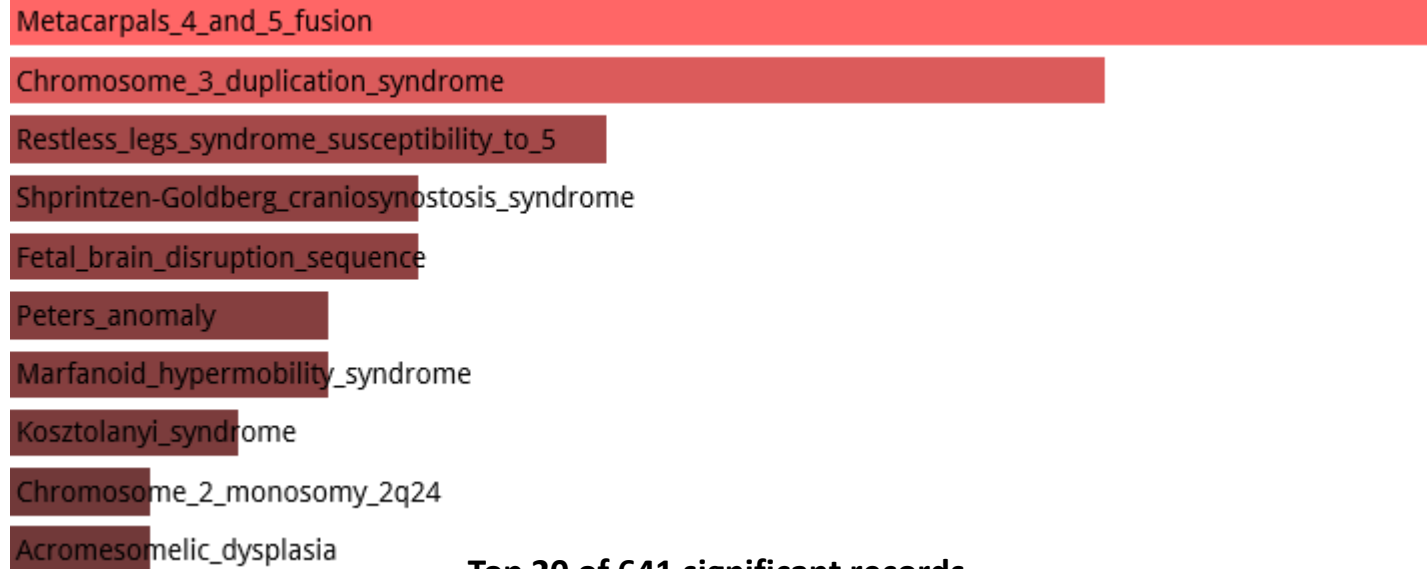


## Enriched Terms

Input Genes



# Rare Diseases AutoRIF ARCHS4 Predictions: 8,405 genes



## Top 20 of 641 significant records

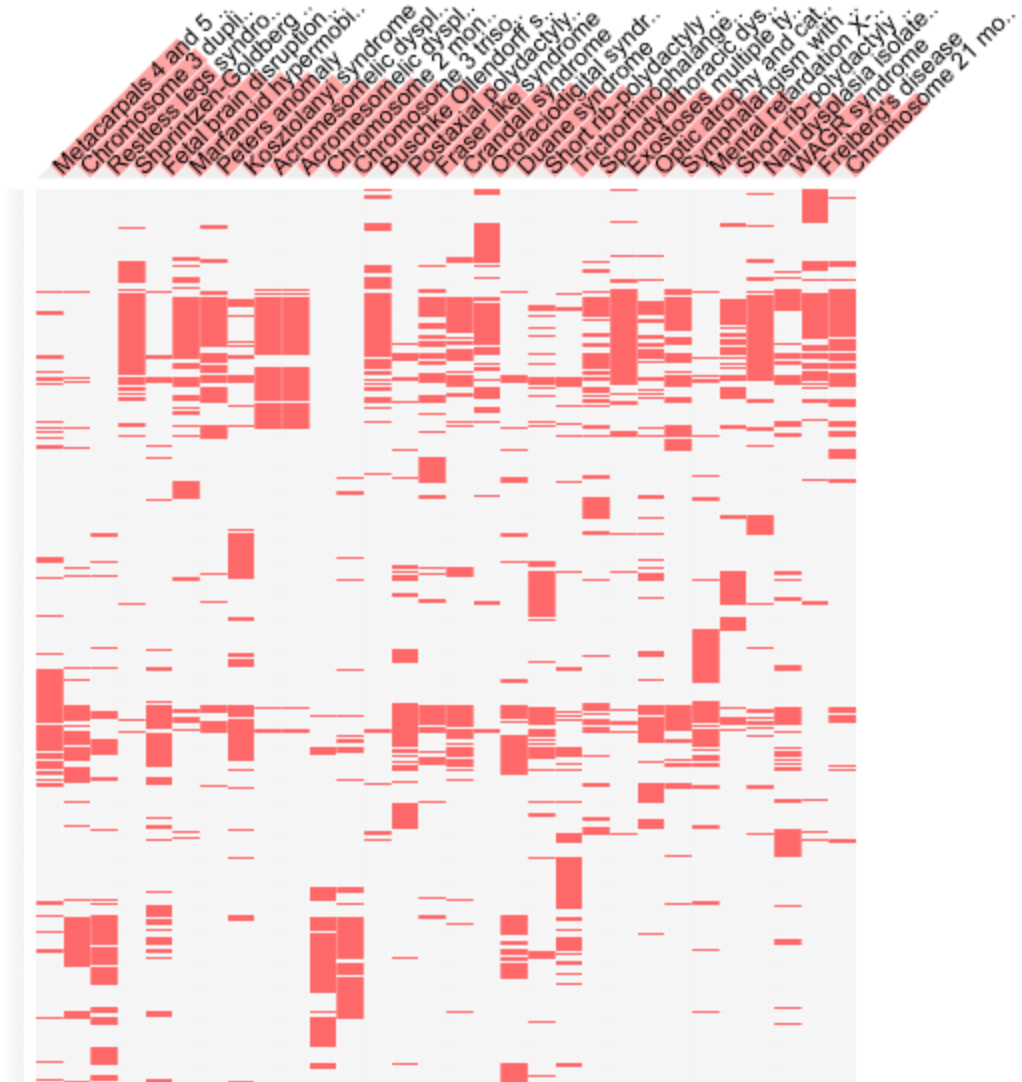
Term	Overlap	P-value	Adjusted P-value
Metacarpals_4_and_5_fusion	163/200	1.33E-30	4.97E-27
Chromosome_3_duplication_syndrome	160/200	2.69E-28	5.01E-25
Restless_legs_syndrome_susceptibility_to_5	155/200	1.01E-24	1.25E-21
Shprintzen-Goldberg_craniosynostosis_syndrome	153/200	2.21E-23	1.65E-20
Fetal_brain_disruption_sequence	153/200	2.21E-23	1.65E-20
Peters_anomaly	152/200	9.95E-23	5.3E-20
Marfanoid_hypermobility_syndrome	152/200	9.95E-23	5.3E-20
Kosztolanyi_syndrome	151/200	4.35E-22	2.03E-19
Chromosome_2_monosomy_2q24	150/200	1.85E-21	6.28E-19
Acromesomelic_dysplasia_Hunter_Thompson_type	150/200	1.85E-21	6.28E-19
Acromesomelic_dysplasia	150/200	1.85E-21	6.28E-19
Chromosome_3_trisomy_3p	149/200	7.69E-21	2.2E-18
Buschke-Ollendorff_syndrome	149/200	7.69E-21	2.2E-18
Fraser_like_syndrome	148/200	3.11E-20	6.81E-18
Crandall_syndrome	148/200	3.11E-20	6.81E-18
Orofaciodigital_syndrome_11	148/200	3.11E-20	6.81E-18
Postaxial_polydactyly_mental_retardation	148/200	3.11E-20	6.81E-18
Short_rib-polydactyly_syndrome_type_4	147/200	1.22E-19	2.28E-17
Trichorhinophalangeal_syndrome_type_3	147/200	1.22E-19	2.28E-17
Duane_syndrome	147/200	1.22E-19	2.28E-17

# Rare Diseases AutoRIF ARCHS4 Predictions: 8,405 genes

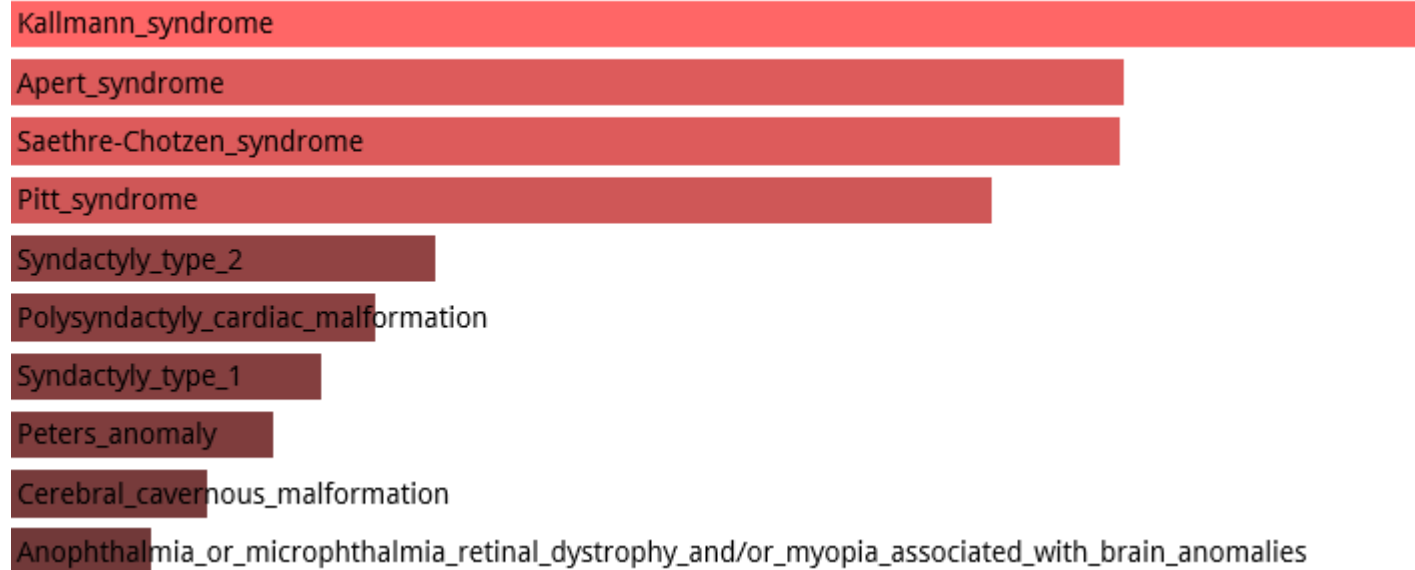


## Enriched Terms

Input Genes



## Rare Diseases AutoRIF Gene Lists: 8,405 genes



### Top 20 of 1,116 significant records

Term	Overlap	P-value	Adjusted P-value
Kallmann_syndrome	159/247	9.25E-13	3.25E-09
Apert_syndrome	143/219	2.58E-12	3.25E-09
Saethre-Chotzen_syndrome	136/206	2.62E-12	3.25E-09
Pitt_syndrome	241/410	4.01E-12	3.73E-09
Syndactyly_type_2	124/188	2.62E-11	1.95E-08
Polysyndactyly_cardiac_malformation	77/104	3.2E-11	1.98E-08
Syndactyly_type_1	122/185	3.85E-11	2.04E-08
Peters_anomaly	75/101	4.53E-11	2.1E-08
Cerebral_cavernous_malformation	157/252	5.65E-11	2.33E-08
Anophthalmia_or_microphthalmia_retinal_dystrophy_and/or_myopia_associated_with_brain_anomalies	182/301	6.82E-11	2.36E-08
X-linked_periventricular_heterotopia	89/126	6.98E-11	2.36E-08
Aniridia	171/282	1.85E-10	5.73E-08
Dominant_cleft_palate	163/267	2.37E-10	6.76E-08
Craniofacial_and_skeletal_defects	94/137	2.64E-10	7.01E-08
Anodontia	140/223	2.84E-10	7.03E-08
Glaucoma_congenital	202/345	3.48E-10	8.08E-08
Childhood-Onset_Schizophrenia	168/278	3.87E-10	8.46E-08
Hennekam_syndrome	139/223	6.68E-10	1.38E-07
Kurczynski_Casperson_syndrome	114/176	8.57E-10	1.68E-07
Osteochondroma	177/299	1.31E-09	2.44E-07



# Rare Diseases AutoRIF Gene Lists: 8,405 genes



## Enriched Terms

Input Genes

