

SUPPLEMENTARY TABLE S4. GENOME COVERAGE SUMMARY OF SEQUENCING DATA

Sample name	Before combining sequencing data				After combining sequencing data			
	Whole exome		X-panel		Whole exome + X-Panel		Whole exome + X-panel (X-chromosome only)	
	Mean coverage (x)	%Bases above 20x	Mean coverage (x)	%Bases above 20x	Mean coverage (x)	%Bases above 20x	Mean coverage (x)	%Bases above 20x
Family1 II.1	55.23	88	350.7	99	73.7	89.8	370.11	99.2
Family1 II.2	59.74	90.3	367.32	99	78.66	91	407.87	99.4
Family1 III.1	55.89	89	448.43	99.3	79.32	89.8	485.46	99.5
Family1 III.2	50.67	86.4	323.64	98.9	67.86	88.5	341.4	99.2
Family1 IV.1	61.1	90.9	468.79	99.3	85.68	91.7	510.16	99.5
Family1 IV.3	52.17	87.4	374.7	99.1	71.98	89.3	392.92	99.3
Family1 IV.2	51.81	87.1	286.34	98.9	66.87	89.1	304.47	99.2
Family1 IV.4	55.33	88.6	361.65	99.1	74.48	90.3	380.98	99.3
Family2 III.1	53.66	86.8	199.01	97.8	63.55	88.4	216.98	98.7
Family2 III.2	50.72	85.2	205.9	97.5	61.02	87	222.92	98.5
Family2 II.1	62.26	89	256.39	98	74.93	90.3	276.78	98.7
Family3 II.1	40.57	78	260.49	98.3	54.07	80.6	274.11	98.8
Family3 II.2	56.1	88.4	340.77	98.6	73.35	89.1	376.48	99.3
Family3 III.1	42.68	79.7	240.05	98.1	54.97	82	254.48	98.7
Family3 III.2	45.63	82	319.53	98.8	62.77	84.6	335.38	99