

### Supplementary Table 3

Association results for previously reported candidate genes for Exfoliation syndrome.

CHR	SNP	BP	A1	F_A	F_U	A2	P	OR	Gene	Reference
1	rs6691985	17289911	G	0.4482	0.4691	A	0.13	0.92	<i>MFAP2</i>	16
1	rs3754511	17310019	G	0.1429	0.1423	A	0.95	1.01	<i>MFAP2</i>	16
1	rs3738815	17314702	G	0.4761	0.4832	A	0.61	0.97	<i>MFAP2</i>	16
1	rs2076604	17318065	A	0.472	0.479	G	0.61	0.97	<i>MFAP2</i>	16
7	rs1015932	147191337	A	0.3256	0.3304	G	0.71	0.98	<i>CNTNAP2</i>	10
7	rs7781317	147201446	G	0.2261	0.2163	A	0.39	1.06	<i>CNTNAP2</i>	10
7	rs971818	147203197	G	0.3406	0.3429	A	0.86	0.99	<i>CNTNAP2</i>	10
7	rs10226360	147206439	A	0.3608	0.3626	C	0.89	0.99	<i>CNTNAP2</i>	10
7	rs12703926	147234727	A	0.2179	0.2342	G	0.16	0.91	<i>CNTNAP2</i>	10
7	rs851821	147242624	G	0.4272	0.4053	A	0.11	1.09	<i>CNTNAP2</i>	10
7	rs11975946	147311353	G	0.3692	0.3866	A	0.19	0.93	<i>CNTNAP2</i>	10
8	rs3087554	27455442	G	0.3364	0.3082	A	0.029	1.14	<i>CLU</i>	16
8	rs11136000	27464519	A	0.2668	0.2863	G	0.11	0.91	<i>CLU</i>	16
9	rs2149356	120474199	A	0.3708	0.3846	C	0.30	0.94	<i>TLR4</i>	17
14	rs2190876	74932268	A	0.2089	0.2492	G	0.00048 <sup>a</sup>	0.80	<i>LTBP2</i>	16
14	rs2302114	74983841	A	0.3935	0.3859	G	0.57	1.03	<i>LTBP2</i>	16
14	rs3815328	74992383	A	0.3763	0.3598	G	0.21	1.07	<i>LTBP2</i>	16
14	rs3784024	75045389	A	0.4215	0.3935	G	0.039	1.12	<i>LTBP2</i>	16
14	rs8014087	75049457	G	0.191	0.1873	A	0.73	1.03	<i>LTBP2</i>	16
14	rs8006778	75071093	A	0.189	0.186	G	0.78	1.02	<i>LTBP2</i>	16
14	rs888414	75104905	A	0.1301	0.1368	G	0.48	0.94	<i>LTBP2</i>	16
14	rs12892228	75108290	A	0.1284	0.1343	G	0.53	0.95	<i>LTBP2</i>	16
14	rs10047892	75111346	A	0.3287	0.345	G	0.21	0.93	<i>LTBP2</i>	16
15	rs1807301	48910240	A	0.4734	0.4971	G	0.085	0.91	<i>FBN1</i>	16
19	rs4803455	41851509	A	0.4276	0.4557	C	0.039	0.89	<i>TGFB1</i>	16
20	rs6023527	36783599	C	0.217	0.2054	A	0.30	1.07	<i>TGM2</i>	16
20	rs4811529	36788191	G	0.3365	0.351	A	0.27	0.94	<i>TGM2</i>	16

a: The initial finding reported by Krumbiegel et al., was in the opposite direction where the A allele conferred susceptibility to XFS.