

## Supplementary Table S2

Predictions of bioinformatics tools for variants segregating in families affected with Bardet-Biedl syndrome (BBS)

Family Identity	Genes	Variants	Provean	Sift	Polyphen2,	Mutation taster
<b>LUBS-01</b> <b>LUBS-02</b> <b>LUBS-09</b>	BBS6/ MKKS	c.775delA, p.Thr259LeuTer2 1	Deleterious	Not applicable	Disease causing	Not applicable
<b>LUBS-03</b>	BBS6/ MKKS	c.748G>A, p.gly250Arg	Deleterious	Damaging	Disease causing	Probably Damaging
<b>LUBS-04</b>	BBS9	c.223C>T, p.Arg75Ter	Deleterious	Not applicable	Disease causing	Not applicable
<b>LUBS-05</b>	BBS1	c.1150 C>T, p.Glu384Ter	Stop gain	Deleterious	Not applicable	Disease causing
<b>LUBS-06</b>	BBS2	c.471 +1G>A	Splicing error	Splicing error	Splicing error	Splicing error
<b>LUBS-10</b>	BBS9	c.252delA, p.Lys85SerTer39	Deleterious	Not applicable	Disease causing	Not applicable
<b>CB-03</b> <b>CB-44</b>	BBS7	c.580delGCA, p.Ala194del	Deleterious	Damaging	Disease causing	Probably Damaging
<b>VI-65</b>	ARL6	c.387_394delAA ATAAAA	Deleterious	Not applicable	Disease causing	Not applicable
<b>RP-04</b>	MKKS	1226G>A, p.Gly409Glu	Deleterious	Damaging	Disease causing	Probably Damaging