

Table S1. Single-SNP association values base on the allele frequencies characterized for the RA patients vs. healthy controls.

| SNP | Position | Gene. Location | A1 | MAF RA | MAF controls | p value | P bonf | OR | L95 | U95 |
|-----------|----------|-------------------------------|----|--------|--------------|-----------------------|-----------------------|------|------|------|
| rs4959028 | 32383138 | BTNL2 HLA-DRA. Intergenic | G | 0.06 | 0.08 | 0.13 | 1.00 | 0.73 | 0.49 | 1.10 |
| rs2157335 | 32386131 | BTNL2 HLA-DRA. Intergenic | A | 0.04 | 0.06 | 0.16 | 1.00 | 0.71 | 0.44 | 1.15 |
| rs9501626 | 32400344 | BTNL2 HLA-DRA. Intergenic | A | 0.15 | 0.19 | 0.02 | 0.49 | 0.73 | 0.55 | 0.96 |
| rs3135335 | 32401845 | BTNL2 HLA-DRA. Intergenic | G | 0.26 | 0.34 | 1.59x10 ⁻³ | 0.03 | 0.70 | 0.55 | 0.87 |
| rs2027856 | 32402705 | BTNL2 HLA-DRA. Intergenic | T | 0.15 | 0.19 | 0.03 | 0.53 | 0.73 | 0.55 | 0.96 |
| rs9268614 | 32402778 | BTNL2 HLA-DRA. Intergenic | G | 0.28 | 0.22 | 5.54x10 ⁻³ | 0.11 | 1.39 | 1.10 | 1.76 |
| rs3129867 | 32404220 | BTNL2 HLA-DRA. Intergenic | G | 0.25 | 0.33 | 7.82x10 ⁻⁴ | 0.02 | 0.68 | 0.54 | 0.85 |
| rs2395178 | 32405362 | BTNL2 HLA-DRA. Intergenic | G | 0.26 | 0.34 | 1.59x10 ⁻³ | 0.03 | 0.70 | 0.55 | 0.87 |
| rs3129882 | 32409530 | HLA-DRA. Intron | G | 0.38 | 0.46 | 2.90x10 ⁻³ | 0.06 | 0.73 | 0.59 | 0.90 |
| rs3129886 | 32410576 | HLA-DRA. Intron | T | 0.18 | 0.22 | 0.06 | 0.22 | 0.78 | 0.60 | 1.01 |
| rs3129888 | 32411726 | HLA-DRA. Intron | G | 0.14 | 0.19 | 0.01 | 0.21 | 0.69 | 0.52 | 0.92 |
| rs2239802 | 32411846 | HLA-DRA. Intron | G | 0.16 | 0.19 | 0.12 | 1.00 | 0.80 | 0.61 | 1.06 |
| rs2395182 | 32413317 | HLA-DRA HLA-DRB1. Intergenic | G | 0.16 | 0.19 | 0.12 | 1.00 | 0.80 | 0.61 | 1.06 |
| rs9268844 | 32429153 | HLA-DRA HLA-DRB1. Intergenic | G | 0.28 | 0.21 | 2.40x10 ⁻³ | 0.05 | 1.44 | 1.14 | 1.83 |
| rs3129768 | 32595083 | HLA-DRB1 HLA-DQA1. Intergenic | G | 0.13 | 0.18 | 0.01 | 1.00 | 0.69 | 0.52 | 0.92 |
| rs7775228 | 32658079 | HLA-DQB1 HLA-DQA2. Intergenic | C | 0.12 | 0.16 | 8.19x10 ⁻³ | 0.16 | 0.66 | 0.49 | 0.90 |
| rs9275224 | 32659878 | HLA-DQB1 HLA-DQA2. Intergenic | A | 0.27 | 0.36 | 2.31x10 ⁻⁴ | 1.61x10 ⁻³ | 0.66 | 0.53 | 0.82 |
| rs9275580 | 32679462 | HLA-DQB1 HLA-DQA2. Intergenic | G | 0.38 | 0.30 | 1.16x10 ⁻³ | 0.02 | 1.43 | 1.15 | 1.77 |
| rs2858332 | 32681161 | HLA-DQB1 HLA-DQA2. Intergenic | G | 0.29 | 0.33 | 0.12 | 1.00 | 0.84 | 0.67 | 1.05 |
| rs3998158 | 32681992 | HLA-DQB1 HLA-DQA2. Intergenic | C | 0.34 | 0.26 | 1.73x10 ⁻³ | 0.03 | 1.43 | 1.14 | 1.78 |

SNP = single nucleotide polymorphism; A1 = minor allele nucleotide; MAF = minor allele frequency; P BONF = *p* value using Bonferroni correction; OR = Odds ratio; L95 = lower bound on confidence interval for odds ratio; U95 = upper bound on confidence interval for odds ratio.

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