

#### S4. Ipathway analysis table

| Pathway name                           | Pathway Id | p-value         | p-value (FDR)   | p-value (Bonferroni) |
|--|------------|-----------------|-----------------|----------------------|
| Olfactory transduction                 | 04740      | <b>7.478e-7</b> | <b>2.311e-4</b> | <b>2.311e-4</b>      |
| Primary immunodeficiency *             | 05340      | <b>7.754e-6</b> | <b>5.245e-4</b> | <b>0.002</b>         |
| Protein digestion and absorption *     | 04974      | <b>7.754e-6</b> | <b>5.245e-4</b> | <b>0.002</b>         |
| Hematopoietic cell lineage *           | 04640      | <b>7.754e-6</b> | <b>5.245e-4</b> | <b>0.002</b>         |
| Cytokine-cytokine receptor interaction | 04080      | <b>8.488e-6</b> | <b>5.245e-4</b> | <b>0.003</b>         |

\* the p-value corresponding to the pathway was computed using only over-representation analysis.