Table S1. Statistical analysis of relative amount of obtained sequences per isotype over the total number of sequences from all 14 donors.

| isotypes | correlation | p-value |
|-----------------|-------------|---------|
| IgA1 | -0.20284 | 0.48675 |
| IgA2 | -0.32861 | 0.25132 |
| IgD | 0.58232 | 0.02889 |
| IgE | 0.24094 | 0.40666 |
| IgG1 | -0.45860 | 0.09908 |
| IgG2 | -0.47966 | 0.08262 |
| IgG3 | -0.43607 | 0.11906 |
| IgG4 | -0.44497 | 0.11086 |
| IgM | 0.70830 | 0.00458 |
| IgM + IgD | 0.70644 | 0.00474 |
| IgA + IgE + IgG | -0.70644 | 0.00474 |

For all nine isotypes and groups correlation and p-values according a linear model fit (F-test) were calculated for age dependency.