

Supplementary Table 1: Description of the SHIP-TREND-0 study sample.

| | | Overall | HLA-typed | |
|---------------------------------|----------------------|-------------|-------------------|------------------|
| | | | DKMS ¹ | WGS ² |
| N | | 976 | 574 | 402 |
| Females | N (%) | 544 (55.7) | 325 (56.6) | 219 (54.5) |
| Males | N (%) | 432 (44.3) | 249 (43.4) | 183 (45.5) |
| Age | mean (SD) | 50.0 (13.7) | 45.3 (14.9) | 56.7 (7.6) |
| | range | 20-81 | 20-81 | 44-70 |
| BMI | mean (SD) | 27.3 (4.6) | 26.8 (4.6) | 28.1 (4.3) |
| Smoking status | | | | |
| Current | N (%) | 214 (21.9) | 158 (27.5) | 56 (13.9) |
| Former | N (%) | 350 (35.8) | 178 (31.0) | 172 (42.8) |
| Never | N (%) | 412 (42.2) | 238 (41.5) | 174 (43.3) |
| <i>S. aureus</i> carrier status | | | | |
| Non-carrier | N (%) | 727 (74.5) | 423 (73.7) | 304 (75.6) |
| Carrier | N (%) | 249 (25.5) | 151 (26.3) | 98 (24.4) |
| Carrier CC30+ | N (% among carriers) | 48 (19.3) | 30 (19.9) | 18 (18.4) |
| Carrier <i>tst</i> + | N (% among carriers) | 49 (19.7) | 30 (19.9) | 19 (19.4) |

¹ DKMS, SHIP-TREND-0 sub-sample that was HLA-typed via PCR-based sequencing by the German Bone Marrow Donor Center (DKMS)

² WGS, SHIP-TREND-0 sub-sample that was HLA-typed based on whole genome sequencing data using the HLA-HD algorithm