



Figure S4. Contribution of each technology to improve the coverage of the single mapping units (SMU) when it performed as the best one. Data are presented for “embryo” (top, 3 technologies; HiSeq, MiSeq and PacBio), “2 cm female cones” (bottom left, 2 technologies: HiSeq and PacBio) and “female cones at time of pollination” (bottom right, 2 technologies: HiSeq and PacBio) samples. For each sample, data are shown by using all size selected or only 3-6 Kb size selected Pacbio libraries.