Figure S7 Recombination rates estimated by LDhat (red) and based on Paigen et al. [S2] (green). As a comparison we also included the Cox et al. [S1] genetic map (blue dashed line). Rates were smoothed over a window of 10 Mb with a shift of 1 Mb. The three maps coincide well. The calculation of the normalized recombination rates is described in Materials and methods.