Flow cytometric analysis of syngeneic tumors. Additional data from the D2A1/D2A1-m2 (n=7-8 mice per group) and 4T07/4T1 (n=6-8 mice per group) tumors analyzed in Fig. 4. A, D2A1/D2A1-m2 and 4T07/4T1 tumor weights at necropsy prior to flow cytometry analysis (mean values ± SEM, unpaired t-test). B, Example of gating strategy employed on a 4T1 tumor sample. C, Quantification of CD4+ FoxP3+ cell (Treg), CD11b+ Ly6G- Ly6C- F4/80+ cell (macrophage) and CD11b+ Ly6C<sub>low</sub> Ly6G+ cell (neutrophil) content of D2A1/D2A1-m2 and 4T07/4T1 tumors (mean values ± SEM, unpaired t-test).