

The scatter plot displays 100 data points, each labeled with a gene name and a category (SUR or DCM). The points are distributed across the plot area, with a clear separation between the SUR and DCM groups. The SUR group is concentrated on the left side of the plot (negative x-values), while the DCM group is concentrated on the right side (positive x-values). A vertical dashed line at x=0 and a horizontal dashed line at y=0 provide a reference for the distribution. The labels for the points include gene names followed by their category, such as 78_SUR, 16_SUR, 82_SUR, 85_SUR, 84_SUR, 85_DCM, 123_SUR, 16_DCM, 45_SUR, 1_DCM, 66_SUR, 30_DCM, 30_SUR, 32_SUR, 18_DCM, 34_SUR, 32_DCM, 34_DCM, 38_DCM, 123_DCM, 82_DCM, 102_DCM, 67_SUR, 68_SUR, 22_SUR, 19_SUR, 24_SUR, 25_SUR, 28_SUR, 70_SUR, 124_SUR, 125_SUR, 126_SUR, 127_SUR, 128_SUR, 129_SUR, 130_SUR, 131_SUR, 132_SUR, 133_SUR, 134_SUR, 135_SUR, 136_SUR, 137_SUR, 138_SUR, 139_SUR, 140_SUR, 141_SUR, 142_SUR, 143_SUR, 144_SUR, 145_SUR, 146_SUR, 147_SUR, 148_SUR, 149_SUR, 150_SUR, 151_SUR, 152_SUR, 153_SUR, 154_SUR, 155_SUR, 156_SUR, 157_SUR, 158_SUR, 159_SUR, 160_SUR, 161_SUR, 162_SUR, 163_SUR, 164_SUR, 165_SUR, 166_SUR, 167_SUR, 168_SUR, 169_SUR, 170_SUR, 171_SUR, 172_SUR, 173_SUR, 174_SUR, 175_SUR, 176_SUR, 177_SUR, 178_SUR, 179_SUR, 180_SUR, 181_SUR, 182_SUR, 183_SUR, 184_SUR, 185_SUR, 186_SUR, 187_SUR, 188_SUR, 189_SUR, 190_SUR, 191_SUR, 192_SUR, 193_SUR, 194_SUR, 195_SUR, 196_SUR, 197_SUR, 198_SUR, 199_SUR, 200_SUR.

Dim 1 (26.92%)