



Figure S15 The genetic affinity of East Asian and Siberian populations to Nganasan and Itelmen measured by Patterson’s $D(\text{Yoruba}, X; \text{Nganasan}, \text{Itelmen})$. In this analysis, we used only the 12 Ainu individuals without non-Ainu ancestry. Both the Sherpa and Tibetans are closer to the Nganasan than to the Itelmen as the other East Asians are. The “WHA” data set used for this analysis includes the Sherpa and Tibetan samples. Horizontal bars around the value represent ± 3 standard deviations.