Normalization strategies differently affect circulating miRNA profile associated with the training status

Martina Faraldi¹*, Marta Gomarasca¹, Veronica Sansoni¹, Silvia Perego¹, Giuseppe Banfi^{1,2}, Giovanni Lombardi¹

- Laboratory of Experimental Biochemistry & Molecular Biology, IRCCS Istituto
 Ortopedico Galeazzi, Milano, Italia
- 2. Vita-Salute San Raffaele University, Milano, Italia

Figure S1. Expression profile of miRNAs ≥ ± 5-fold up- or down-regulated in plasma of sedentary and trained subjects. Expression profile of miRNAs associated to skeletal muscle and bone, which function was defined in experimental models rather than in humans. Statistical analysis was performed with Prism v6.01 (GraphPad Software). All data are expressed as mean ±SEM and data were compared throughout t-test with Welch's correction. The differences were considered significant when p value < 0.05 (*p<0.05, **p<0.01, ***p<0.001).

