Table S7. Candidate genes in the top 1% of PBS results in Collas.

| **Ranka** | **Gene** | **Name** | **Function** | **Hypoxia/ arsenic association** | **Window PBS MAX:****Gene PBS MAX** |
| --- | --- | --- | --- | --- | --- |
| 2 | *CBS* | Serine sulfhydrase  | Bradykinin receptor involved in NO mediated vasodilation; associated with CBF | Cellular response to hypoxia | 1.005:1.005 |
| 52 | *PRKG1* | Protein kinase, cGMP-dependent | Key mediator NO pathway, regulates platelet activation and adhesion, smooth muscle contraction, cardiac function | NO stimulates guanylate cyclase | 0.693:0.693 |
| 137 | *STC2* | Stanniocalcin 2 | Bone and skeletal muscle growth; HIF-1 activated, protects cells from apoptosis in hypoxia | Cellular response to hypoxia | 0.583:41 kb downstream0.050 |
| 172 | *FOXO1* | Forkhead box protein O1A  | Transcription factor: main target of insulin signalling; regulates metabolic homeostasis in response to oxidative stress | Cellular response to ROS | 0.558:0.558 |
| 190 | *UBE2D3* | Ubiquitin-protein ligase D3 | E2 ubiquitin-conjugating enzyme, targets p53 and EGFR among others | Cellular response to hypoxia | 0.548:0.548 |
| 198 | *SOD1* | Superoxide dismutase 1  | Convert superoxide radicals to molecular oxygen and hydrogen peroxide | Cellular response to ROS | 0.545:38 kb downstream0.208 |
| 202 | *GNGT1* | Guanine nucleotide-binding protein G(T1)  | GMPase, mediates activation by rhodopsin of a cyclic GTP-specific phosphodiesterase | Cellular response to hypoxia | 0.543:0.543 |

aLower ranks are reported for PBS as window size was halved compared to other tests.