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| **EKM Gauss D=20 λ=0.3** - Gauss SVM (C=10, σ2=1) accuracy 93.57% (**20 genes**) |
| 878: ACCNUM=M87789; gene 1 IG GAMMA-1 CHAIN C REGION (HUMAN);  **306**: ACCNUM=J00231; gene 1 Human Ig gamma3 heavy chain disease OMM protein mRNA;  2: ACCNUM=R39465; 3' UTR 2a 23933 EUKARYOTIC INITIATION FACTOR 4A (Oryctolagus cuniculus);  3: ACCNUM=????;  11: ACCNUM=T72863; 3' UTR 1 84277 FERRITIN LIGHT CHAIN (HUMAN);  10: ACCNUM=T55131; 3' UTR 1 73931 "GLYCERALDEHYDE 3-PHOSPHATE DEHYDROGENASE, LIVER (HUMAN)  6: ACCNUM= R02593; 3' UTR 2a 124094 60S ACIDIC RIBOSOMAL PROTEIN P1 (Polyorchis penicillatus)  24: ACCNUM= T59954; 3' UTR 1 79441 THYMOSIN BETA-4 (HUMAN)  5: ACCNUM= U14973; gene 1 "Human ribosomal protein S29 mRNA, complete cds.  15: ACCNUM= U14971; gene 1 "Human ribosomal protein S9 mRNA, complete cds.  26: ACCNUM= T95018; 3' UTR 2a 120032 40S RIBOSOMAL PROTEIN S18 (Homo sapiens)  8: ACCNUM= H80240; 3' UTR 2a 240814 INTER-ALPHA-TRYPSIN INHIBITOR COMPLEX COMPONENT II PRECURSOR (Homo sapiens)  17: ACCNUM= L28809; gene 1 "Homo sapiens dbpB-like protein mRNA, complete cds.  22: ACCNUM= J02763; gene 1 "Human calcyclin gene, complete cds  33: ACCNUM=T51574; 3' UTR 1 72258 40S RIBOSOMAL PROTEIN S24 (HUMAN)  119: ACCNUM=T72175; 3' UTR 1 85528 IG KAPPA CHAIN PRECURSOR V-III REGION (HUMAN)  47: ACCNUM=T58861; 3' UTR 2a 77563 60S RIBOSOMAL PROTEIN L30E (Kluyveromyces lactis)  31: ACCNUM=T61609; 3' UTR 1 78081 LAMININ RECEPTOR (HUMAN);  13: ACCNUM=X63432; gene 1 H.sapiens ACTB mRNA for mutant beta-actin (beta'-actin  44: ACCNUM=T88723; 3' UTR 1 109876 UBIQUITIN (HUMAN) |