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S.Pounds, C.Cheng, X.Cao, K.R.Crews, W.Plunkett, V.Gandhi, J.Rubinstein, B.Richter, J.D.Downing and J.Lamb

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C.Lu and R.D.King

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K.F.Kerr

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APPLICATIONS NOTE

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PROCAIN server for remote protein sequence similarity search
Y.Wang, R.I.Sedravy and N.V.Gishin

The Sequence Alignment/Map format and SAMtools
H.Li, B.Handaker, A.Wysoker, T.Fennell, J.Ruan, N.Homer, G.Marth, G.Abecasis, D.Defost and 1000 Genome Project Data Processing Subgroup

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