**International Pea Sequencing Consortium Go-to-Meeting**

**6 March 2012**

**Conference call proposed agenda**:

1. Introductions

        Principles of cooperation-discussion

        Ultimate goal of the consortium-discussion

 Please read the attached notes for this discussion:

2. Obtain a survey sequence for pea genome (genome zipper, Mayer et al 2009; 2001, attached)

        Is this an approach to pursue?

        Which pea line?  Cameor suggested, BAC resource available.

        Summary of public pea sequences/transcriptomes available

        a. Conduct survey (‘shotgun’) sequencing

        b. Synteny analysis

 3.  Develop SNP markers

        Are more needed?

        50 genotypes, which ones?

 4.  Sequence the genome (‘Gold standard’)

        Again, which pea line?

        Physical map

        Anchoring the physical map

        Sequencing

5. Funding sources