# **Supplementary information**

# Yeast-based bioproduction of disulfide-rich peptides and their cyclization via asparaginyl endopeptidases

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# Yeast-based bioproduction of cyclic disulfide-rich peptides and their cyclization via asparaginyl endopeptidases

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### pPICZ-alpha-SFTI-KLK5

MRFPSIFTAVLFAASSALAAPVNTTTEDETAQIPAEAVIGYSDLEGDFDVAVLPFSNSTNNGLLFINTT IASIAAKEEGVSLEKR<mark>GFCHRSYPPECWPNGLPHHHHHHH</mark>

### pPICZ-alpha-[G22N] cVc1.1

MRFPSIFTAVLFAASSALAAPVNTTTEDETAQIPAEAVIGYSDLEGDFDVAVLPFSNSTNNGLLFINTT IASIAAKEEGVSLEKRGCCSDPRCNYDHPEICGGAAGNGLPHHHHHHHH

### pPICZ-alpha-MCoTI-II

MRFPSIFTAVLFAASSALAAPVNTTTEDETAQIPAEAVIGYSDLEGDFDVAVLPFSNSTNNGLLFINTT IASIAAKEEGVSLEKRGGVCYGRRARRRCRRDSDCPGACICKGNGYCGSGSDGLPHHHHHHHH

### b



### pET-15-ubiquitin-[C247A]OaAEP1<sub>b</sub>

MHHHHHSSGLVPRGSHMQIFVKTLTGKTITLEVEPSDTIENVKAKIQDKEGIPPDQQRLIFAGKQLED GRTLSDYNIQKESTLHLVLRLRGGARDGDYLHLPSEVSRFFRPQETNDDHGEDSVGTRWAVLIAGSKGY ANYRHQAGVCHAYQILKRGGLKDENIVVFMYDDIAYNESNPRPGVIINSPHGSDVYAGVPKDYTGEEVN AKNFLAAILGNKSAITGGSGKVVDSGPNDHIFIYYTDHGAAGVIGMPSKPYLYADELNDALKKKHASGT YKSLVFYLEACESGSMFEGILPEDLNIYALTSTNTTESSWAYYCPAQENPPPPEYNVCLGDLFSVAWLE DSDVQNSWYETLNQQYHHVDKRISHASHATQYGNLKLGEEGLFVYMGSNPANDNYTSLDGNALTPSSIV VNQRDADLLHLWEKFRKAPEGSARKEVAQTQIFKAMSHRVHIDSSIKLIGKLLFGIEKCTEILNAVRPA GQPLVDDWACLRSLVGTFETHCGSLSEYGMRHTRTIANICNAGISEEQMAEAASQACASIP

### pET-15-ubiquitin-MCoAEP2

MHHHHHSSGLVPRGSHMQIFVKTLTGKTITLEVEPSDTIENVKAKIQDKEGIPPDQQRLIFAGKQLED GRTLSDYNIQKESTLHLVLRLRGGDLPGDFLRLPSEALKFFHRGASDATGDEDSVGTRWAVLIAGSNGY WNYRHQADICHAYQLLRKGGLKDENIIVFMYDDIAFNPENPRPGVIINHPHGSDVYHGVPKDYTGEDVN VENFFAAILGDKKAIKGGSGKVVDSGPNDHIFIFYSDHGGPGVLGMPTYPYIYADGLIDVLKKKHASGS YKSLVFYLEACESGSIFEGLLPQDLNIYATTASNAVESSWGCYCPGDDTAPPPEYDTCLGDLYSVGWME DSDRHNLKTESLRQQYELVKKRTLNDYTVYGSHVMQYGDITLNKNALFSYLGTDPANENNTFVESNSLR PTTKVTNQRDADMVHFWEKFRKAPEGSAQKIEAQKHFVEAMSHRVHIDNSVKLIGKLLFGIEKGPEVLN AVRPTGQPLVNNWDCLKNMVRSFETHCGSLSQYGMKHMRSFANLCNAGIRNEQMAEASAQACVSVPSGP WSSLHKGFTA

Supplementary Fig. 1 Overview of construct design for CPs and recombinant AEPs. a The amino acid sequences of *S. cerevisiae*  $\alpha$ -mating factor secretion signal (grey colored letters) fused on the N-terminus of cyclic peptide precursors (red colored letters) followed by a C-terminus His tag (blue colored letters). **b** The amino acid sequences of AEP zymogens represented (green colored letters) fused to the C-terminus of ubiquitin solubility protein (grey colored letters) and a His tag was placed on the N-terminus of the ubiquitin protein (blue colored letters).



**Supplementary Fig. 2 Bioreactor production of MCoTI-II CPs. a** Parameters of a bioreactor bioproduction of MCoTI-II precursors showing temperature, dissolved oxgen (DO) and pH for the growth phases of glycerol batch, glycerol-fed batch and methanol fed batch. **b** SDS-PAGE analysis of MCoTI-II CP accumulation at different time point post methanol induction. Expected bands indicated by a magenta colored arrow.



Supplementary Fig. 3 HPLC-UV-HRMS chromatograms. a IMAC purified fractions containing MCoTI-II precursor before AEP-mediated cyclization. b RP-HPLC purified recombinantly produced MCoTI-II. <sup>&</sup> unidentified *P. pastoris* contaminant and  $^{\beta}$  - aspartyl isomers<sup>1</sup>.

1 Hernandez, J. F. *et al.* Squash trypsin inhibitors from Momordica cochinchinensis exhibit an atypical macrocyclic structure. *Biochemistry* **39**, 5722-5730 (2000).