1	Supplementary material
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4	Statistical Evaluation to Validate Matrix-Matched Calibration for
5	Standardized Beany Odor Compounds Quantitation in Yellow Pea Flour
6	Using HS-SPME-GC-MS
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17 1. Figures and tables

18 Fig. S1 (A) Soxhlet extraction, (B) Accelerated dichloromethane extraction (ASEDCM),
19 Raffinate from (C1) three-cycle of accelerated dichloromethane extraction of starch & (C2)
20 three-cycle of accelerated dichloromethane extraction of yellow pea, (D) Deodorized matrix-

- 21 matched yellow pea flour from accelerated dichloromethane extraction (ASEDCMYP), and
- 22 (E) Native yellow pea flour.





25 Fig. S2 External standard calibration curves of six beany odor makers established in methanol

26 via weighted least square regression





1-6 denote as beany odor markers 1-hexanal, 3-methyl-1-butanol, 1-hexanol, (3Z)-hexen-1-ol, 1-nonanal, and 2-

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ethyl-1-hexanol, respectively.

- 30 Fig. S3 Residual plots of six beany odor makers in two deodorized matrix-matched flours via
- 31 ordinary least square (OLS) and weighted least square (WLS) regression.

Litomotumo			С	ompound				Quantification		
reported	1-hexanal	3-methyl-1- butanol 1-hexa		(3Z)-hexen- 1-ol	1-nonanal	2-ethyl-1- hexanol	Pea details	Processing	methods	
1	$\checkmark$	$\checkmark$	$\checkmark$	$\checkmark$	$\checkmark$		Commercial green pea (GP)	Unblanched GP	No	
2		$\checkmark$	$\checkmark$	$\checkmark$			GP grown at Richmond, N.S.W.	Blanched GP	No	
3	$\checkmark$	$\checkmark$	$\checkmark$	$\checkmark$	$\checkmark$		GP variety <i>medullare</i> Alef in Denmark	Blanched GP	Semi-quantification with IS	
4	$\checkmark$						6 YP, 3 GP cultivars in Saskatchewan	Crop years 2005-2007	Ratio of relative peak area	
5	$\checkmark$	$\checkmark$	$\checkmark$	$\checkmark$	$\checkmark$		Commercial pea flour	SAFE/SPME extraction	Semi-quantification with IS	
6	$\checkmark$	$\checkmark$	$\checkmark$	$\checkmark$	$\checkmark$		Commercial YP	Germination from 1 to 6 days	Calibration curves in deodorized starch	
7	$\checkmark$		$\checkmark$		$\checkmark$		YP Zhong-wan variety	Crop year 2017, raw and roasted YP	Calibration curves in deodorized YP flour	
8	$\checkmark$	$\checkmark$	$\checkmark$		$\checkmark$	$\checkmark$	YP Agassize and Trapeze cultiar	Crop year 2018	Ratio of relative peak area	
9	$\checkmark$	$\checkmark$	$\checkmark$	$\checkmark$	$\checkmark$	$\checkmark$	Commercial YP	Different milling configurations	Calibration curves in deodorized YP flour	

Table S1. Literature investigation on beany odor quantification in pea cultivars

 $34 \quad \sqrt{\text{ indicates the compound has been reported in literature.}}$ 

35 GP: green pea, YP: yellow pea, IS: internal standard, SAFE: solvent assisted flavor evaporation; SPME: solid phase microextraction.

			Concentration of	each beany	odor standard in mi	xed SS/ mg/n	nL	
Compound		1-hexanal	3-methyl-1-butanol	1-hexanol	(3Z)-hexen-1-ol	1-nonanal	2-ethyl-1-butanol	
Weight/g		0.1063	0.0977	0.1076	0.1020	0.1045	0.1134	
Meth	anol/mL	10	10	10	10	10	10	
<b>SS</b> <sup>1</sup> /	mg/mL	10.63	9.77	10.76	10.2	10.45	11.34	
Code	DF <sup>2</sup>	(	Concentration of each b	eany odor sta	andard in diluted mi	ixed solutions	s / μg/mL	
B25	2.50×10 <sup>3</sup>	0.45	0.41	2.27	0.43	0.22	0.48	
B50	5.00×10 <sup>3</sup>	0.22	0.21	1.13	0.22	0.11	0.24	
D	$1.00 \times 10^{4}$	0.11	0.10	0.57	0.11	0.06	0.12	
A50	1.25×10 <sup>4</sup>	0.09	0.08	0.45	0.08	0.04	0.10	
C25	5 $2.50 \times 10^4$ 0.05 0.1		0.04	0.23	0.04	0.02	0.05	
C50	$5.00 \times 10^{4}$	0.02	0.02	0.11	0.02	0.01	0.02	
B2525	6.25×10 <sup>4</sup>	0.018	0.016	0.091	0.017	0.009	0.019	
Е	$1.00 \times 10^{5}$	0.011	0.010	0.057	0.011	0.006	0.012	
B5025	1.25×10 <sup>5</sup>	0.009	0.008	0.045	0.009	0.004	0.010	
D25	2.50×10 <sup>5</sup>	0.004	0.004	0.023	0.004	0.002	0.005	
D50	5.00×10 <sup>5</sup>	0.002	0.002	0.011	0.002	0.001	0.002	
F	$1.00 \times 10^{6}$	0.001	0.001	0.006	0.001	0.0006	0.001	
C5025	$1.25 \times 10^{6}$	0.0009	0.0008	0.0045	0.0009	0.0004	0.0010	
E25	$2.50 \times 10^{6}$	0.0005	0.0004	0.0023	0.0004	0.0002	0.0005	
E50	$1.25 \times 10^{8}$	0.0002	0.0002	0.0011	0.0002	0.0001	0.0002	

37 Table S2. Preparation of beany odor mixed stock solutions and corresponding dilution factors

38 1. SS represents stock solution in methanol.

39 2. DF represents dilution factor, diluted with methanol.



The codes and concentrations are corresponding to those in Table S2.

## 44 2. Material and methods

45 2.3 Physical attributes of deodorized matrix-matched flours

46 2.3.1 Surface morphology

47 Surface morphology of deodorized matrix-matched flour from different treatment 48 combinations were observed via JEOL JSM-6300 Scanning Electron Microscope (SEM, JEOL 49 USA, Peabody, MA USA)<sup>9</sup>. The flour samples were adhered to aluminum mounts by carbon 50 tabs and coated with gold using a Balzers SCD 030 sputter coater (BAL-TEC RMC, Tucson, 51 AZ USA). Images were taken at × 1000 magnification with an accelerating voltage of 15 kV.

52 2.3.2 Particle size distribution

Particle size distribution (PSD) of deodorized matrix-matched flour from different treatment combinations were determined by a laser light scattering instrument (Mastersizer 3000, Malvern Instruments Ltd., Malvern, U.K.) in dry powder sampling module<sup>9</sup>. The particle size was determined from intensity-time fluctuations of the laser light scattered from a sample at a fixed angle of 173°. Measurements were reported as the volume-weight mean diameter (D  $_{[4,3]} = \sum n_i d_i^4 / \sum n_i d_i^3$ ), where  $n_i$  was the number of particles of diameter  $d_i$ .

59 2.4 Beany odor analysis

60 2.4.1 beany odor detection

The beany odor detection and identification were achieved by headspace-solid phase microextraction-gas chromatography-mass spectrometry (HS-SPME-GC-MS, 7890B GC system, 5977A mass detector, Agilent Technologies, Santa Clara, CA, USA.) under scanning and selective ion mode (SIM) according to previous study with minor modifications<sup>9</sup>.

65 A SPME fiber coated with DVB/CAR/PDMS (50/30 µm, Supelco Inc. Bellefonte, PA) was applied to extract the volatile compounds. PAL RSI 120 autosampler (CTC Analytics, 66 Zwingen, Switzerland) was employed for the incubation and extraction procedure. An aliquot 67 of 0.5 g YP flour mixed with 2 mL of deionized water was transferred in a 20 mL glass vial 68 capped by aluminum caps with PTFE/silicone septa (diam.  $\times$  thickness 20 mm  $\times$  0.105 in, 69 MilliporeSigma, St. Louis, MO, USA). Vials were incubated for 5 min at 60 °C in an 70 autosampler heating block, while agitating at 250 rpm. Thereafter, the SPME fiber needle was 71 inserted into the vial for 60 min to adsorb volatiles at 60 °C, which was then transferred into 72 73 the injector port (250 °C) for 3 min desorption.

74 The separation of volatiles was performed on an Agilent Technologies 7890B GC system. The injection port was operated in splitless mode. The liner was Agilent 5190-2293: 900 µL 75 76 (Splitless, single taper, Ultra Inert) and the purge flow to split vent was 50 mL/min at 1 min. Volatiles were separated on a J&W DB-Wax column (60 m × 0.25 mm i.d., 0.25 µm film 77 thickness, Agilent Technologies, Santa Clara, CA, USA.). Oven temperature program was set 78 as: initial temperature 40 °C, followed by an increase of 45 °C/min to 85 °C, then by 3 °C/min 79 to 135 °C and held for 1 min, next by 1 °C/min to 160 °C and held for 1 min, afterwards by 9 80 °C/min to 200 °C, and lastly by 45 °C/min to 250 °C and held for 12 min. Helium carrier gas 81 (purity 99.9999%, Praxair, Fargo, ND, USA.) was used at a flow rate of 2 mL/min. The column 82 83 effluent was transferred to the 5977A mass detector, operated in electron impact (EI) ionization mode at 70 eV. The temperature of MS detector transfer line was hold at 250 °C. The MS 84 quadrupole temperature was 150 °C. Ion source temperature was 230 °C. Scanning and 85 selecting ion monitoring mode (SIM) was applied with scan time and segments setting from 4 86

87 to 63 min and scanning m/z from 40 to 350.

### 88 2.4.2 Identification of beany odor markers

89 The beany odor markers were identified by three methods: i) raw data acquired by GC-MS were analyzed using Agilent MassHunter Qualitative Analysis (B.07.00) coupled with MS 90 91 library NIST 17. The min match score > 85 was the main criteria applied for validating the entities; ii) to calculate the retention index (RI) of each compound using a series of n-alkanes 92 C8-C40 (MilliporeSigma, St. Louis, MO, USA). Calculated RI was matched with a reference 93 value according to NIST database (https://webbook.nist.gov); and iii) standard chemical 94 95 compounds were applied to further verify the compounds under the aforementioned GC-MS conditions. 96

The retention time (RT) around 9.5 min with a SIM quantifier ion of 56.1 was assigned to 1-hexanal (9.5 min-56.1  $m/_z$ , **1**). The remaining beany odor makers were: 15.1 min-55.1  $m/_z$  (**2**), 15.3 min-43  $m/_z$  (**3**), 16.1 min-41  $m/_z$  (**4**), 16.6 min-56.1  $m/_z$  (**5**), 19.6 min-57.1  $m/_z$  (**6**), and 12.3 min-85  $m/_z$  (**IS**), respectively. The detailed information regarding SIM ions parameters is listed

101 below.

Time/mi n	Group name	Number of ions	Mass1/dwell time/s	Mass2/dwell time/s	Mass3/dwell time/s
9.5	Hexanal	3	41.00/50	44.10/50	56.10*/50
12.3	2-Methyl-3-heptanone	3	57.00/50	85.00*/50	128.00/50
15.1	3-Methyl-1-butanol	3	42.00/50	55.10*/50	70.10/50
15.3	1-Hexanol	3	43.00/50	55.10/50	56.10/50
16.1	(3Z)-Hexen-1-ol	3	41.00*/50	67.00/50	82.10/50
16.6	Nonanal	3	41.00*/50	56.10/50	57.10/50

	19.6	2-Methyl-1-hexanol	3	41.00/50	43.00/50	57.10*/50
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102 \* represented quantitative SIM ion.

## 103 3. Results and discussion

#### 104 3.4 Visual Evaluation of Linearity of Calibration Curves

As displayed in Fig. S3, there were strong evidence of heteroskedasticity in the response of each beany odor marker via OLSR in both deodorized matrix-matched flours as the distribution was neither placed randomly alongside zero, nor shown an increasing behavior of the deviations. Similar residual plot results were also reported by Gomes et al. in pesticides guantitation by ESC method<sup>10</sup>.

110 Outliers were vastly detected in compound 1, 2, 4, 5, 6 in ASEDCMS2D matrix, whereas they were only noticed in compound 2 and 6 in ASEDCMYP2D matrix. After removing the 111 112 outliers, the studentized residual by WLSR showed a more discrete and random distribution, especially in ASEDCMYP2D matrix. Take compound 3 as an example, the residual 113 distribution was completely dispersed alongside the concentration, and each replicate was 114 almost discrete from each other in ASEDCMYP2D matrix when regressed by WLSR. 115 Conversely, it showed a declined trend and later fanned out in ASEDCMYP2D matrix with 116 117 OLSR. Besides that, there seemed to be an interesting phenomenon that the residual 118 distribution patterns of each beany odor marker were more random in ASEDCMYP2D matrix than in ASEDCMS2D matrix with the same WLSR, in which certain distribution patterns of 119 120 the latter still showed a bit fan-out trend such as compound 5.

- 121 3.6 Beany Odor Compounds Quantitation
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A <sub>i</sub> <sup>56.1</sup>	$A_{IS}{}^{85} \\$	$A_{i}^{56.1}\!/A_{IS}^{85}$	${m_i}^{56.1}\!/{m_{IS}}^{85}$	mi <sup>56.1</sup> -ASEDCMYP/ug	m <sup>56.1</sup> -ASEDCMYP/ug/g	Mean	STD
2520.64	259.74	9.70	0.23	0.37	0.75	1 11	0.51
3193.58	188.80	16.92	0.46	0.74	1.47	1.11	0.51

125 Taking hexanal in ASEDCMYP2D as an example,

 $A_i^{56.1}$  was the response peak area of SIM ion 56.10f hexanal reading from the TIC;

128 A<sub>IS</sub><sup>85</sup> was the response peak area of SIM ion 85 of internal standard reading from the TIC;

 $m_i^{56.1}/m_{IS}^{85}$  was interpolated according to the calibration curve  $A_i^{56.1}/A_{IS}^{85} = 31.648 m_i^{56.1}/m_{IS}^{85} + 2.331$ ;

130 m<sub>i</sub><sup>56.1</sup>-ASEDCMYP/ug was achieved by multiple the mass of added IS, which was 1.598 μg;

 $m_i^{56.1}$ -ASEDCMYP/ug/g was achieved by dividing the applied sample mass of 0.5 g.

132 The final quantitative results were reported by mean  $\pm$  std.

## 133 Taking hexanal in ASEDCMS2D as an example,

A <sub>i</sub> <sup>56.1</sup>	$A_{IS}{}^{85} \\$	$A_i{}^{56.1}\!/A_{IS}{}^{85}$	${m_i}^{56.1}\!/{m_{IS}}^{85}$	mi <sup>56.1</sup> -ASEDCMS/ug	mi <sup>56.1</sup> -ASEDCMS/ug/g	Mean	STD
2697.71	495.20	5.45	0.50	0.79	1.59	1.47	0.17
2575.21	526.4	4.892	0.42	0.67	1.35		

 $m_i^{56.1}/m_{IS}^{85}$  was interpolated according to the calibration curve  $A_i^{56.1}/A_{IS}^{85} = 7.159 m_i^{56.1}/m_{IS}^{85} + 1.872;$ 

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