

Module	Process	Factor	Description	Default Value	Unit
phenology	thermal time accumulation	x_temp	Cardinal temperatures affecting the effective thermal time (y_tt = 0 26 0 oCd)	0 26 34	°C
phenology	germination failure (time)	days_germ_limit	Maximum days allowed after sowing for germination to take place	40	days
phenology	coleoptile elongation (water)	fasw_emerg	Fraction of available soil water affecting the stress factor [0-1] for thermal time calculation between germination and emergence (rel_emerg_rate = 1 1)	0 1	[0,1]
phenology	emergence (temperature)	shoot_lag	Time lag before linear coleoptile growth starts	40	°Cd
phenology	coleoptile elongation (temperature)	shoot_rate	Degree days required to increase coleoptile by 1 mm	1.5	°Cd mm-1
phenology	emergence failure (temperature)	tt_emerg_limit	Maximum degree days allowed for emergence to take place	300	°Cd
phenology	node phyllochron	y_node_app_rate	Thermal time between the appearance of successive nodes as a function of the node number of the main stem (x_node_no_app = 0 20)	95 95	°Cd
phenology	leaf initiation	y_leaves_per_node	Leaf per node defined as a function of x_node_no_leaf=0 2.5 6	1 1 6	leaf node-1
phenology	node development (carbon)	y_leaf_no_frac	Leaf appearance depending on LAI (x_lai_ratio = 0 0.1 0.8 1)	0 1 1	[0-1]
phenology	phenology failure	swdf_pheno_limit	Critical cumulative phenology water stress, above which the crop fails	25	-
phenology	development failure (temperature)	x_weighted_temp	Temperature affecting plant death (y_plant_death = 0 1)	55 60	°C
cultivar	phenological stages shift (photoperiod)	photop_sens	Sensitivity to photoperiod	3	-
cultivar	phenological stages shift (vernalisation)	vern_sens	Sensitivity to vernalisation	1.5	-
cultivar	thermal time accumulation (juvenile)	tt_end_of_juvenile	Thermal time between plant emergence and end of juvenile stage	400	°Cd
cultivar	thermal time accumulation (floral initiation)	tt_floral_initiation	Thermal time between end of juvenile and floral initiation	555	°Cd
cultivar	thermal time accumulation (flowering)	tt_flowering	Thermal time between floral initiation and flowering	120	°Cd
cultivar	thermal time accumulation (grain filling)	tt_start_grain_fill	Thermal time between flowering and start of grain filling	545	°Cd
cultivar	stem elongation	y_height	plant canopy height for stem weight per plant (x_stem_wt = 0 6 g plant-1)	0 1500	mm
cultivar	grain development	grains_per_gram_stem	Number of grains that are set depending on the stem dry weight	25	grain g-1
cultivar	grain demand (flowering)	potential_grain_growth_rate	Potential rate of grain growth from flowering to grain fill	0.001	g d-1
cultivar	grain demand (maturity)	potential_grain_filling_rate	Potential rate of grain growth during grain filling	0.002	g d-1
cultivar	grain demand (maximum)	max_grain_size	Potential grain size	0.041	g
biomass	initial leaf biomass	leaf_dm_init	Initial leaf dry matter	0.003	g
biomass	initial root biomass	root_dm_init	Initial root dry matter	0.01	g
biomass	initial stem biomass	stem_dm_init	Initial stem dry matter	0.0016	g
biomass	photosynthesis (co2)	co2_rue_modifier	Factor affecting the CO2 compensation point	163	-
biomass	biomass growth (generic temperature)	tfac_slope	Temperature stress factor slope	0.0125	-
biomass	biomass growth (generic temperature, min)	temp_fac_min	Temperature stress factor minimum	0.69	-
biomass	biomass growth (generic water)	sfac_slope	Soil water stress factor slope	-0.125	-
biomass	biomass growth (generic water, max)	sw_fac_max	Soil water stress factor maximum	1.125	-
biomass	biomass growth (temperature)	x_ave_temp	Daily mean temperatures affecting the RUE multiplier [0-1] (y_stress_photo = 0 1 1 0)	0 10 25 35	°C
biomass	radiation interception (green)	y_extinct_coef	Extinction coefficient of green leaves as a response to row spacing (x_row_spacing = 200 350 1000 mm)	0.5 0.5 0.5	-
biomass	radiation interception (dead)	y_extinct_coef_dead	Extinction coefficient for dead leaves as a response to row width (x_row_spacing = 200 350 1000 mm)	0.3 0.3 0.3	-
biomass	grain demand (temperature)	x_temp_grainfill	Temperatures affecting stress index [0-1] for the potential grain filling rate (y_rel_grainfill = 0 1 1)	0 26 35	°C
biomass	grain demand (nitrogen)	x_temp_grain_n_fill	Temperatures affecting stress index [0-1] for potential N filling rate (y_rel_grain_n_fill = 0 1 1)	0 25	°C
biomass	photosynthesis (potential)	y_rue	Radiation Use Efficiency depending on stages	0 0 1.24 1.24 1.24 1.24 1.24 1.24 0 0 0	g MJ-1
biomass	root biomass partitioning	y_ratio_root_shoot	Root/shoot ratio as a function of specific stages (x_stage_no_partition = 1 2 3 4 4.9 5 5.4 6 6.9 7 8 9 10 11)	0 0 1 1 0.3 0.3 0.3 0.08 0.01 0 0 0 0 0	[0,1]
biomass	leaf biomass partitioning	y_frac_leaf	Fraction of remaining dry matter allocated to leaves for specific stages (specific stages (x_stage_no_partition = 1 2 3 4 4.9 5 5.4 6 6.9 7 8 9 10 11)	0 0 0.6 0.6 0.6 0.42 0 0 0 0 0 0 0	[0,1]
biomass	pod biomass partitioning	y_frac_pod	Fraction of dry matter allocated to rachis for specific stages (x_stage_no_partition = 1 2 3 4 4.9 5 5.4 6 6.9 7 8 9 10 11)	0 0 0 0 0 0.33 0.33 0.33 0 0 0 0 0	[0,1]
biomass	grain water content	grn_water_cont	Water content of grain	0.125	g g-1

Module	Process	Factor	Description	Default Value	Unit
leaf	leaf expansion (init)	initial_tpla	Initial total plant leaf area	200	mm ²
leaf	leaf expansion (carbon)	y_sla	Parameter added to change minimum and maximum values all at once	1	-
leaf	leaf expansion	node_no_correction	Number of growing leaves in the sheath	2	leaf
leaf	leaf expansion (carbon)*	sla_min	Minimum specific leaf area	18000	mm ² mg ⁻¹
leaf	leaf expansion (carbon)*	y_sla_max	Maximum specific leaf area as a function of daily change in LAI (x_lai = _sla_max)	27000 22000	mm ² g ⁻¹
leaf	leaf expansion (failure)	leaf_no_crit	Critical number of leaves, below which portion of the crop may die due to water stress	10	leaf
leaf	leaf expansion (potential)	y_leaf_size	Leaf potential size depending on the number of nodes appeared (x_node_no = 1 5 8 10) and the growing leaf number in the sheath (node_no_correction = 2)	1400 3700 4800 5600	mm ²
root	root absorption (capacity)	ll_modifier	Parameter added to change crop lower limit values for all depths at once	1	-
root	root absorption (potential)	kl_modifier	Parameter added to change crop kl values for all depths at once	1	-
root	root elongation (phenology)	root_depth_rate	Potential rate of root front increase for the different stages	0 5 30 30 30 30 0 0 0 0	mm day ⁻¹
root	root elongation (soil structure)	xf_modifier	Parameter added to change crop xf values for all depths at once	1	-
root	root elongation (temperature)	x_temp_root_advance	Temperatures affecting root front by the [0-1] stress index: y_rel_root_advance = 0 1 0	0 25 35	°C
root	root elongation (carbon)	specific_root_length	Specific root length	105000	mm g ⁻¹
root	root elongation (carbon)	y_rel_root_rate	Factor affecting root length density in a layer, which depends on root branching (x_plant_rld = 0 0.0001 0.0003 mm/mm ³ /plant)	0.02 0.1 1	[0,1]
root	root elongation (generic water)	x_sw_ratio	Water availability affecting the stress factor for root depth growth (y_sw_fac_root = 0 1)	0 0.25	[0,1]
root	root elongation (init)	initial_root_depth	Initial depth of roots	100	mm
root	root elongation (water)	x_ws_root	Water supply demand ratio affecting the increase in root depth (y_ws_root_fac = 1 1)	0 1	[0,1]
water	phenological stages shift (water)	x_sw_avail_ratio	Water availability that affects stress indices for phenology: y_swdef_pheno = 1.0 1.0	0 0.5	[0,1]
water	leaf expansion (water)	x_sw_demand_ratio	Water availability that affects stress indices for leaf expansion: y_swdef_leaf = 0 1	0.1 1.1	[0,1]
water	photosynthesis failure (water)	swdf_photo_limit	Critical cumulative photosynthesis water stress, above which the crop partly fails	99	-
water	transpiration efficiency (phenology)	transp_eff_cf	Transpiration efficiency coefficient for aboveground biomass for the different stage codes	0 0 0.006 0.006 0.006 0.006 0.006	g m ⁻² mm ⁻¹
water	water demand (VPD)	eo_crop_factor_default	Default crop factor for limiting soil	1.5	kPa
water	transpiration efficiency (co2)	y_co2_te_modifier	CO ₂ factor for transpiration efficiency in response to x_co2_te_modifier = 350 700 1000 ppm (small hypothesis)	1 1.69	-
nitrogen	initial leaf nitrogen	leaf_n_init_conc	Leaf initial N concentration	0.063	g g ⁻¹
nitrogen	initial root nitrogen	root_n_init_conc	Root initial N concentration	0.02	g g ⁻¹
nitrogen	initial stem nitrogen	stem_n_init_conc	Stem initial N concentration	0.05	g g ⁻¹
nitrogen	initial pod nitrogen	pod_n_init_conc	Rachis initial N concentration	0.063	g g ⁻¹
nitrogen	nitrogen supply	kno3	Constant for extractable soil nitrogen	0.02	-
nitrogen	nitrogen supply (potential)	total_n_uptake_max	Maximum total nitrogen uptake in a soil layer	0.6	g m ⁻²
nitrogen	phenological stages shift (nitrogen)	N_fact_pheno	Multiplier for N deficit effect on phenology	100	-
nitrogen	leaf expansion (nitrogen)	N_fact_expansion	Multiplier for N deficit effect on leaf expansion	1	-
nitrogen	photosynthesis (nitrogen)	N_fact_photo	Multiplier for nitrogen deficit effect on photosynthesis	1.5	-
nitrogen	leaf nitrogen remobilisation	leaf_n_sen_conc	N concentration that remains in senesced leaves (i.e. that is not retranslocated)	0.005	g g ⁻¹
nitrogen	grain filling (nitrogen)	N_fact_grain	Multiplier for N deficit effect on grain filling	1.5	-
nitrogen	leaf nitrogen demand (min, crit, max)	n_conc_leaf	Parameter added to change minimum, critical and maximum values of n_conc_leaf all at once	1	-
nitrogen	leaf nitrogen demand*	y_n_conc_crit_leaf	Critical leaf N concentration at the different stages	0.063 0.063 0.042 0.015 0.008 0.0035	g g ⁻¹
nitrogen	leaf nitrogen demand*	y_n_conc_max_leaf	Maximum leaf N concentration at the different stages	0.07 0.07 0.05 0.02 0.01 0.005	g g ⁻¹
nitrogen	leaf nitrogen demand*	y_n_conc_min_leaf	Minimum leaf N concentration at the different stages	0.025 0.025 0.021 0.01 0.005 0.0025	g g ⁻¹
nitrogen	root nitrogen demand (min, crit, max)	n_conc_root	Parameter added to change minimum, critical and maximum values of n_conc_root all at once	1	-
nitrogen	root nitrogen demand*	n_conc_crit_root	Critical root N concentration at the different stages	0.02	g g ⁻¹
nitrogen	root nitrogen demand*	n_conc_max_root	Maximum root N concentration at the different stages	0.02	g g ⁻¹
nitrogen	root nitrogen demand*	n_conc_min_root	Minimum root N concentration at the different stages	0.01	g g ⁻¹
nitrogen	root nitrogen remobilisation	root_n_sen_conc	N concentration that remains in the senesced roots (i.e. that is not retranslocated)	0.005	g g ⁻¹
nitrogen	stem nitrogen demand (min, crit, max)	n_conc_stem	Parameter added to change minimum, critical and maximum values of n_conc_stem all at once	1	-
nitrogen	stem nitrogen demand*	y_n_conc_crit_stem	Critical stem N concentration at the different stages	0.05 0.05 0.02 0.01 0.005 0.0035	g g ⁻¹
nitrogen	stem nitrogen demand*	y_n_conc_max_stem	Maximum stem N concentration at the different stages	0.07 0.07 0.04 0.015 0.015 0.015	g g ⁻¹
nitrogen	stem nitrogen demand*	y_n_conc_min_stem	Minimum stem N concentration at the different stages	0.025 0.025 0.004 0.003 0.003 0.0025	g g ⁻¹
nitrogen	stem nitrogen remobilisation	stem_n_sen_conc	N concentration that remains in the senesced stems (i.e. that is not retranslocated)	0.0025	g g ⁻¹

Module	Process	Factor	Description	Default Value	Unit
nitrogen	pod nitrogen demand (min, crit, max)	n_conc_pod	Parameter added to change minimum, critical and maximum values of n_conc_pod all at once	1	-
nitrogen	pod nitrogen demand*	y_n_conc_crit_pod	Critical rachis N concentration at the different stages	0.05 0.05 0.02 0.01 0.005 0.0035	g g-1
nitrogen	pod nitrogen demand*	y_n_conc_max_pod	Maximum rachis N concentration at the different stages	0.07 0.07 0.04 0.015 0.015 0.015	g g-1
nitrogen	pod nitrogen demand*	y_n_conc_min_pod	Minimum rachis N concentration at the different stages	0.025 0.025 0.004 0.003 0.003 0.0025	g g-1
nitrogen	pod nitrogen remobilisation	pod_n_sen_conc	N concentration that remains in the senesced rachis (spike without grains)	0.007	g g-1
nitrogen	meal nitrogen demand (min, crit, max)	n_conc_meal	Parameter added to change minimum, critical and maximum values of n_conc_meal all at once	1	-
nitrogen	meal nitrogen demand*	y_n_conc_crit_meal	Critical meal N concentration at the different stages	0 0 0 0.03 0.03	g g-1
nitrogen	meal nitrogen demand*	y_n_conc_max_meal	Maximum meal N concentration at the different stages	0 0 0 0.03 0.03	g g-1
nitrogen	meal nitrogen demand*	y_n_conc_min_meal	Minimum meal N concentration at the different stages	0 0 0 0.014 0.014	g g-1
nitrogen	meal nitrogen remobilisation	meal_n_sen_conc	N concentration that remains in senesced grains (i.e. that is not retranslocated)	0.014	g g-1
nitrogen	gain nitrogen demand (min, crit, max)	n_conc_grain	Parameter added to change minimum, critical and maximum values of n_conc_grain all at once	1	-
nitrogen	grain nitrogen demand*	n_conc_crit_grain	Critical grain N concentration at the different stages	0.03	g g-1
nitrogen	grain nitrogen demand*	n_conc_max_grain	Maximum grain N concentration at the different stages	0.03	g g-1
nitrogen	grain nitrogen demand*	n_conc_min_grain	Minimum grain N concentration at the different stages	0.014	g g-1
senescence	leaf senescence (age)	fr_lf_sen_rate	Fraction of total leaf number senescing per main stem node	0.035	[0-1]
senescence	leaf senescence (age)	node_sen_rate	Rate of node senescence on main stem	60	°Cd node-1
senescence	leaf senescence (temperature)	x_maxt_senescence	Minimum temperature affecting the daily fraction of leaf area senescing due to heat (y_heatsenescence_fac = 0 0.05 0.40)	34 34.1 45	°C
senescence	leaf senescence (water)	sen_rate_water	Slope in linear equation relating soil water stress during photosynthesis to leaf senescence rate	0.1	-
senescence	leaf senescence (carbon)	min_tpla	Lower limit of total leaf area per plant	5	mm2 plant-1
senescence	leaf senescence (shading)	lai_sen_light	Critical LAI when shading is starting to cause leaf area senescence	7	m2 m-2
senescence	leaf senescence (shading)	sen_light_slope	Sensitivity of leaf area senescence to shading	0.002	-
senescence	root senescence (nitrogen)	y_dm_sen_frac_root	Rate of root biomass senescing each day as a function of x_dm_sen_frac_root = 0 1	0.005 0.005	[0-1]

Table S1. Description of the APSIM-Wheat parameters included in the sensitivity analysis. *Module* refers to the sub-model where the parameter is used in APSIM-Wheat, *Process* refers to the physiological process targeted by the considered parameter and *Factor* is the parameter name used in the present study and in the APSIM-Wheat documentation (Zheng et al., 2014), where a complete description of the parameters is given. The *Default Value* field lists the nominal value of the parameter for cultivar *Hartog* in APSIM-Wheat 7.5 (only first three values were presented when the parameter is defined as a vector). In the *Process* field, influential parameters in indicated in bold and parameters that were grouped together for physiologic reasons are identified by (*).