SUPPLEMENTAL MATERIAL

Supplementary Material 1 – Heterogeneity of individual kinetic profiles

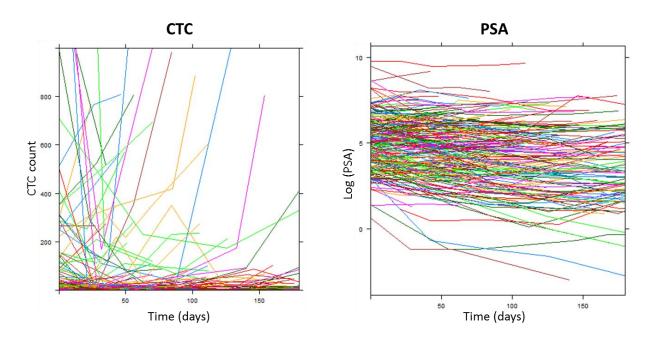
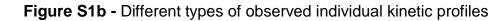
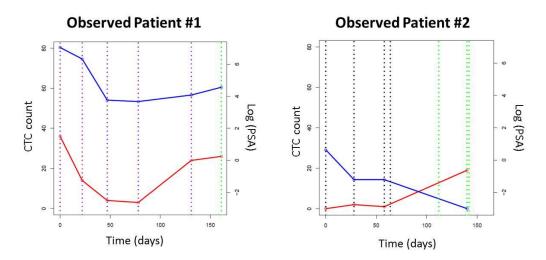


Figure S1a – Spaghetti plots

All 223 individual profiles were plotted for PSA and CTC. Each color corresponds to one individual.





Blue curves represent the PSA kinetics and red curves the CTC kinetics. Vertical dashed lines represent the treatment cycles: chemotherapy in black, hormonotherapy in green, and both treatments administered simultaneously in purple.

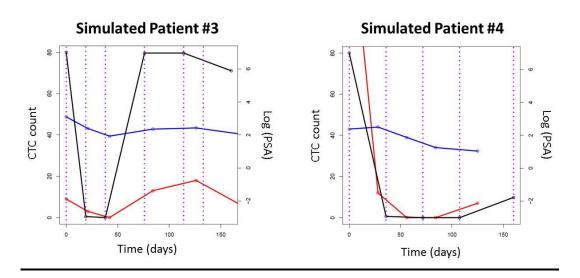
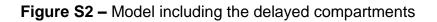


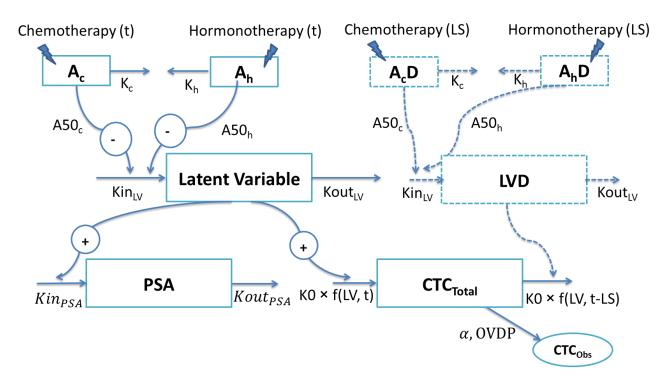
Figure S1c - Different types of simulated individual kinetic profiles

Blue curves represent PSA kinetics, red curves CTC kinetics, and black curves the latent variable kinetics. Vertical dashed lines represent the treatment cycles.

Supplementary Material 2 – Model including the delayed compartments

To estimate the CTC lifespan, the model was implemented in NONMEM using delayed compartments with the ALAG function. Figure S2 represents the model including the delayed compartments.





This model was described by the following equations:

$$\begin{aligned} \frac{dA_c}{dt} &= -K_c \times A_c \\ \frac{dA_cD}{dt} &= -K_c \times A_cD \\ \frac{dA_h}{dt} &= -K_h \times A_h \\ \frac{dA_hD}{dt} &= -K_h \times A_hD \\ \frac{dLV}{dt} &= Kin_{LV} \times \left(1 - \frac{A_c}{A50_c + A_c}\right) \times \left(1 - \frac{A_h}{A50_h + A_h}\right) - Kout_{LV} \times LV \\ \frac{dLVD}{dt} &= Kin_{LV} \times \left(1 - \frac{A_cD}{A50_c + A_cD}\right) \times \left(1 - \frac{A_hD}{A50_h + A_hD}\right) - Kout_{LV} \times LVD \quad (if \ t > LS) \\ \frac{dLVD}{dt} &= 0 \quad (if \ 0 < t < LS) \\ \frac{dPSA}{dt} &= Kin_{PSA} \times LV - Kout_{PSA} \times PSA \\ \frac{dCTC_{Total}}{dt} &= K0 \times LV - K0 \times LVD \end{aligned}$$

 A_c and A_h represent drug amounts in the chemotherapy and the hormonotherapy compartments (*Arbitrary Unit, AU*), respectively. A_cD and A_hD represent drug amounts in the delayed chemotherapy and the delayed hormonotherapy compartments (*Arbitrary Unit, AU*), respectively. K_c and K_h are the chemotherapy and hormonotherapy kinetics rate constants, respectively (day^{-1}). A50_c and A50_h respectively are the amounts of each treatment producing 50% of the maximum effect (*AU*). LV corresponds to the latent variable (*AU*), and Kin_{LV} and Kout_{LV} are their production and elimination rates (AU.day⁻¹ and day⁻¹), respectively. LVD corresponds to the delayed latent variable (*AU*). LVD is supposed to be constant and equal to LV0 for times before the lifespan. Kin_{PSA} and Kout_{PSA} correspond to the PSA production and elimination rates (ng.ml⁻¹.AU⁻¹ and day⁻¹), respectively. K0 is the CTC production rate (CTC.day⁻¹.AU⁻¹).

The initial conditions of the model at time 0 were as follows:

$$\begin{cases} A_{c}(0) = 0\\ A_{c}D(0) = 0\\ A_{h}(0) = 0\\ A_{h}D(0) = 0\\ LV(0) = LV_{0} \text{ with } LV_{0} < \frac{Kin_{LV}}{Kout_{LV}}\\ LVD(0) = LV_{0}\\ PSA(0) = PSA_{0}\\ CTC_{Total}(0) = K0 \times LS \times LV_{0} \end{cases}$$

$$LS = ALAG_{A_cD} = ALAG_{A_hD} = ALAG_{LVD}$$

 LV_0 and PSA_0 are the initial latent variable value (AU) and the initial PSA concentration (ng/ml), respectively. LS correspond to the CTC lifespan (day).

Supplementary Material 3 – Data file and NONMEM code

Fragment of data file:

ID	TIME	AMT	DV	СМТ	MDV
1	0	1		1	1
1	0	1		2	1
1	0	1		5	1
1	0	1		6	1
1	0	1		4	1
1	0		11	4	0
1	0		4.127134	8	0
1	21	1		1	1
1	21	1		2	1
1	21	1		5	1
1	21	1		6	1
1	21		0	4	0
1	21		3.688879	8	0
1	42	1		1	1
1	42	1		2	1
1	42	1		5	1
1	42	1		6	1
1	42		0	4	0
1	42		3.391147	8	0

CMT: 1=Chemotherapy; 2=Hormonotherapy; 4=CTC; 5=Delayed Chemotherapy; 6=Delayed Hormonotherapy; 8=PSA.

NONMEM Code:

\$PROBLEM Joint model of PSA and CTC count kinetics in mCRPC patients **\$INPUT** ID TIME AMT DV CMT MDV **\$DATA** ...

\$SUBROUTINE ADVAN13 TOL=9

\$MODEL NCOMP=8

- COMP = (A1) ; PK chemotherapy
- COMP = (A2) ; PK hormonotherapy
- COMP = (A3) ; Latent Variable
- COMP = (A4); CTC
- COMP = (A5) ; Delayed PK chemotherapy
- COMP = (A6); Delayed PK hormonotherapy
- COMP = (A7) ; Delayed Latent Variable
- COMP = (A8) ; PSA

\$PK CALLFL=-2 ; Call the PK subroutine with every event record, with additional and lagged doses
MU_1=LOG(THETA(1))
LV0=EXP(MU_1+ETA(1)) ; Initial Latent Variable

MU_2=LOG(THETA(2)) Kc=EXP(MU_2+ETA(2))

; Chemotherapy kinetic rate constant

MU 3=LOG(THETA(3)) Kh=EXP(MU_3+ETA(3)) ; Hormonotherapy kinetic rate constant MU_4=LOG(THETA(4)) $A50c=EXP(MU_4+ETA(4))$; Amount of chemotherapy producing 50% of the maximum effect MU 5=LOG(THETA(5)) A50h=EXP(MU_5+ETA(5)) ; Amount of hormonotherapy producing 50% of the maximum effect MU 6=LOG(THETA(6)) KOUTLV=EXP(MU_6+ETA(6)) ; Latent Variable elimination rate constant MU 7=THETA(7) SFLV=EXP(MU_7+ETA(7)) KINLV=(TS0*KOUTTS)/((SFLV)/(1+(SFLV))) ; Latent Variable production rate MU 8=LOG(THETA(8)) KINPSA=EXP(MU 8+ETA(8)) ; PSA production rate MU 9=LOG(THETA(9)) KOUTPSA=EXP(MU_9+ETA(9)) ; PSA elimination rate constant MU_10=LOG(THETA(10)) PSA0=EXP(MU_10+ETA(10)) ; Initial PSA concentration MU_11=THETA(11) K0=MU 11+ETA(11) ; CTC production rate MU 12=THETA(12) ; Delay duration, lifespan (LS) ALAG5=MU 12+ETA(12) ALAG6=ALAG5 ALAG7=ALAG5 F4=K0*ALAG5 ; Initial condition for CTC: CTC(0)=K0*LS*LV0 (LV0=1) MU 13=LOG(THETA(13)) OVDP=EXP(MU_13+ETA(13)) ; Overdispersion MU 14=LOG(THETA(14)) $W1=EXP(MU_14+ETA(14))$: Standard deviation for PSA residual error ; Initial Conditions at time 0: $A_0(1)=0$ A_0(2)=0 A_0(3)=LV0 $A_0(5)=0$ A 0(6)=0 A 0(7)=LV0 A_0(8)=PSA0 **\$DES** $DADT(1) = -Kc^*A(1)$: Time course of chemotherapy amount $DADT(2) = -Kh^*A(2)$; Time course of hormonotherapy amount DADT(3)=KINLV*(1-(A(1)/(A50c+A(1))))*(1-(A(2)/(A50h+A(2))))-KOUTLV*A(3) ; Time course of LV $DADT(5) = -Kc^*A(5)$; Delayed time course of chemotherapy amount $DADT(6) = -Kh^*A(6)$; Delayed time course of hormonotherapy amount DADT(7)=KINLV*(1-(A(5)/(A50c+A(5))))*(1-(A(6)/(A50h+A(6))))-KOUTLV*A(7) ; Delayed time course of LV

A7=LV0 IF(T.GT.ALAG5) A7=A(7)

DADT(4)=K0*A(3)-K0*A(7) ; Time course of total CTCs DADT(8)=KINPSA*A(3)-KOUTPSA*A(8) ; Time course of PSA

\$ERROR

LAMB=A(4)*0.0015 ; Expected number of CTCs in the aliquot

PSA=A(8) nCTC=DV

; Factorial: LFAC=GAMLN(nCTC+1.)

;gamma functions of the negative binomial model expression: LGAM1=GAMLN(nCTC+1/OVDP) LGAM2=GAMLN(1/OVDP) LTRM1=(LOG(1/(1+OVDP*LAMB)))*(1/OVDP) LTRM2=(LOG(LAMB/(LAMB+1/OVDP)))*(nCTC)

;Logarithm of the Negative Binomial distribution: LNB=LGAM1-LFAC-LGAM2+LTRM1+LTRM2

; CTC observation: IF (CMT.EQ.4) THEN F_FLAG=2 Y=-2*LNB ;-2 Log Likelihood ENDIF

; PSA observation: IF (CMT.EQ.8) THEN F_FLAG=0 IPRED=LOG(PSA) Y=IPRED+W1*ERR(1) IRES=DV-IPRED IWRES=IRES/W1 ENDIF

\$THETA

; LV0
; Kc
; Kh
; A50c
; A50h
; KOUTLV
; SFLV
; KINPSA
; KOUTPSA
; PSA0
; K0
; ALAG5
; OVDP
; W1

\$OMEGA

0 FIX ; LV0

\$OMEGA BLOCK(9)

0.5 ; Kc 0.01 0.5 ; Kh 0.01 0.01 0.5 ; A50c 0.01 0.01 0.01 0.5 ; A50h 0.01 0.01 0.01 0.01 0.5 ; KOUTLV 0.01 0.01 0.01 0.01 0.01 0.5 ; SFLV 0.01 0.01 0.01 0.01 0.01 0.01 0.5 ; KINPSA 0.01 0.01 0.01 0.01 0.01 0.01 0.01 0.5

; KOUTPSA ; PSA0

\$OMEGA BLOCK(2)

10 ; K0 1 10 ; ALAG5

\$OMEGA

; OVDP 0.5 0 FIX ;W1

\$SIGMA

1 FIX

\$ESTIMATION METHOD=SAEM LAPLACE INTER NUMERICAL

<u>Supplementary Material 4</u> – Categorical VPCs for the dichotomized CTC count (<5 vs \ge 5)

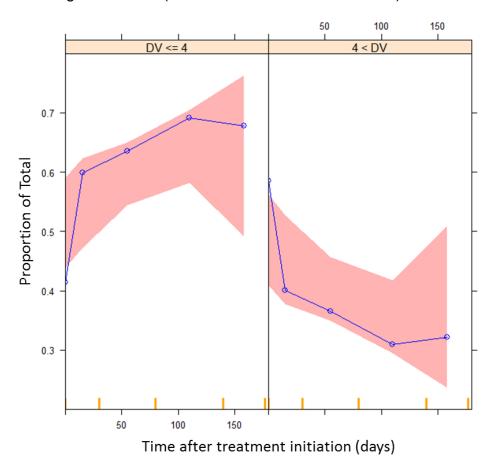


Figure S3 – Categorical VPCs (CTC count <5 vs CTC count \ge 5)

The probabilities of having a number of CTCs lower than 5 or greater than 5 were plotted versus time. Red areas are the 95% confidence intervals of the simulated median probabilities. Blue lines are the observed probabilities.