

# COMPOSITIONS

\* Components  $\{ N_2 \quad CO_2 \quad H_2S \quad H_2O \quad Hg \dots \}$  non-HC

\* Amounts  $(m_i, n_i)$

$C_1 \quad C_2 \quad C_3 \quad i-C_4 \quad n-C_4 \quad i-C_5 \quad n-C_5$

$C_6 \quad C_7 \quad C_8 \quad C_9 \quad C_{10} \quad C_{11} \dots C_{N-1} \quad \boxed{C_{N+}}$

Many Isomers (PNA) reported by normal (latm) boiling points

SCN

Heavy Plus (Residue)

SCN  $C_7$

$n-C_6$  (1st isomer in SCN " $C_7$ ")

Benzene:  $C_6$  Aromatic

$n-C_7$  (highest NBP in " $C_7$ ")

SCN  $C_8$

$n-C_8$

Field A B C

$C_{10}$   $\left\{ \begin{array}{l} GC \\ TBP \end{array} \right.$

SCN overlap:  $\boxed{C_n}$  Distillation

$C_{n-2} \quad C_{n-1} \quad C_n \quad C_{n+1} \quad C_{n+2}$

<5% <15% >60% <15% <5%

	$\frac{C_{N+}}$							
< 1980	$C_{7+}$							} Gamma Molar Distribution Model
1980s	$C_{10+}$	$C_{12+}$	$C_{15+}$					
1990s	$C_{20+}$			w/ $C_6 \quad C_7 \quad C_8 \quad C_9$ isomers				
> 2000	$C_{26+}$	$C_{31+}$	$C_{35+}$	-u-				

Example  
 COMPONENT SLATES  
 JK1 sample

Component
Hydrogen Sulfide
Nitrogen
Carbon Dioxide
Methane
Ethane
Propane
Iso-butane
N-butane
2-2 Dimethylpropane
Iso-pentane
N-pentane
2-2 Dimethylbutane
Cyclopentane
2-3 Dimethylbutane
2 Methylpentane
3 Methylpentane
Other Hexanes
n-Hexane
Heptanes Plus
TOTAL

Component
Hydrogen Sulfide
Nitrogen
Carbon Dioxide
Methane
Ethane
Propane
Iso-butane
N-butane
2-2 Dimethylpropane
Iso-pentane
N-pentane
2-2 Dimethylbutane
Cyclopentane
2-3 Dimethylbutane
2 Methylpentane
3 Methylpentane
Other Hexanes
n-Hexane
Methylcyclopentane
Benzene
Cyclohexane
2-Methylhexane
3-Methylhexane
2,2,4 Trimethylpentane
Other Heptanes - 7
n-Heptane
Methylcyclohexane - 7
Toluene - 7
Other C-8's
n-Octane
Ethylbenzene
M&P-Xylene
O-Xylene
Other C-9's
n-Nonane
Other C10's
n-Decane
Undecanes Plus
TOTAL

Component
Hydrogen Sulfide
Nitrogen
Carbon Dioxide
Methane
Ethane
Propane
Iso-butane
N-butane
2-2 Dimethylpropane
Iso-pentane
N-pentane
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2,2,4 Trimethylpentane
Other Heptanes
n-Heptane
Methylcyclohexane
Toluene
Other C-8's
n-Octane
Ethylbenzene
M&P-Xylene
O-Xylene
Other C-9's
n-Nonane
Other C10's
n-Decane
Undecanes
Dodecanes
Tridecanes

C<sub>6</sub>

C<sub>7</sub>

C<sub>8</sub>

C<sub>9</sub>

C<sub>10</sub>

C<sub>10</sub>  
 C<sub>11</sub>  
 C<sub>12</sub>

Tetradecanes
Pentadecanes
Hexadecanes
Heptadecanes
Octadecanes
Nonadecanes
Eicosanes
Heneicosanes
Docosanes
Tricosanes
Tetracosanes
Pentacosanes
Hexacosanes
Heptacosanes
Octacosanes
Nonacosanes
Triacontanes
Hentriacontanes Plus

C<sub>30</sub>  
 C<sub>31+</sub>

# \* Amounts

lab ⇒ Mass Measures

$$m_i \quad \overline{w}_i = \frac{m_i}{m_t} \quad m_t = \sum_j m_j$$

Moles

$$n_i \quad \left. \begin{matrix} z_i \\ y_i \\ x_i \end{matrix} \right\} = \frac{n_i}{n_t} \quad n_t = \sum_j n_j$$

$$m_i \leftrightarrow n_i$$

$$M_i = \frac{m_i}{n_i}$$

uncertain number for C<sub>6</sub>...

Molar Mass

Molecular Weight

PNA

$$M_i = 14i + h$$

$$h = +2$$

alkanes (paraffins) "P"

$$h = 0$$

naphthene

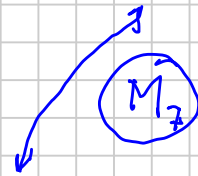
$$h = -6$$

aromatic

Backcalc  
for C<sub>7</sub>  
-20

erg. Benzene C<sub>6</sub>H<sub>6</sub> :  $M_B = 6 \cdot 14 - 6 = \underline{78}$

point of "C<sub>7</sub>" SCN



$$n-C_7 : M = 14 \cdot 7 + 2 = \underline{100}$$

+2

$$KF \text{ (Table 5.2)} = 96$$

"World Average"  
2

# \* Measure Composition Amounts?

## Gas Chromatography (GC)

- (1) For sep. gases & flashed gas
  - (2) For flashed (STO) oils
- } ⇒ GC  $w_{gi} \quad y_i$   
 $w_{oi}$
- flashed at 1 atm

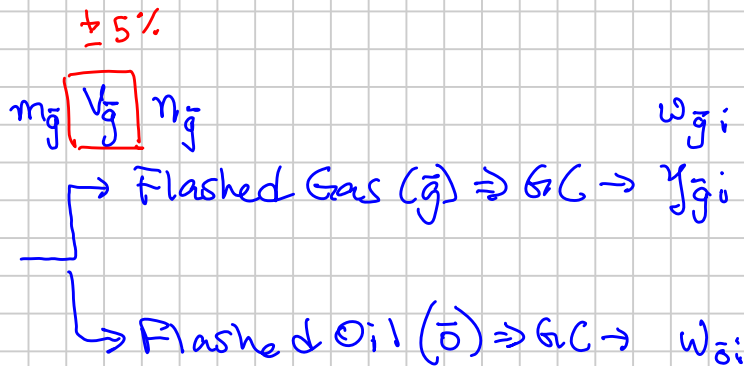
$\bar{g}$  : flash (atm) gas

$\bar{o}$  : flash (atm) oil  
eg. STO

## General Method to Measure Pressurized Sample Compositions

- BHS
- MDT/RCI
- Sep. Oil

① Bring to 1 atm  
@ Ambient



$$GOR_{recomb} = \frac{V_{\bar{g}}}{V_{\bar{o}}}$$

1 atm =  $p, T(t)$

$\pm$   $m_{\bar{o}} \{ \sum_{\bar{o}} M_{\bar{o}} \}$  measured ⇒  $V_{\bar{o}} n_{\bar{o}}$

Recombine mathematically

Original Sample

$$m_i = (m_{\bar{g}} w_{gi} + m_{\bar{o}} w_{oi})$$

$$w_i = \frac{m_i}{m_t}$$

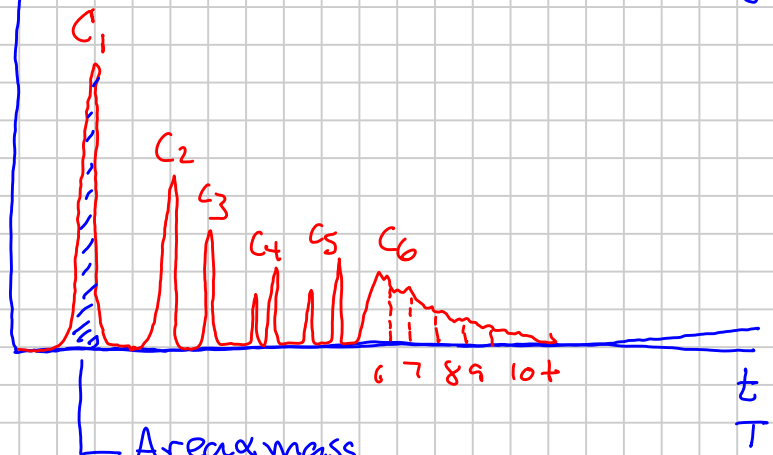
## FLASH-GC-RECOMBINE (Alt. Cryogenic Flash-GC-Rec)

# Gas Chromatography

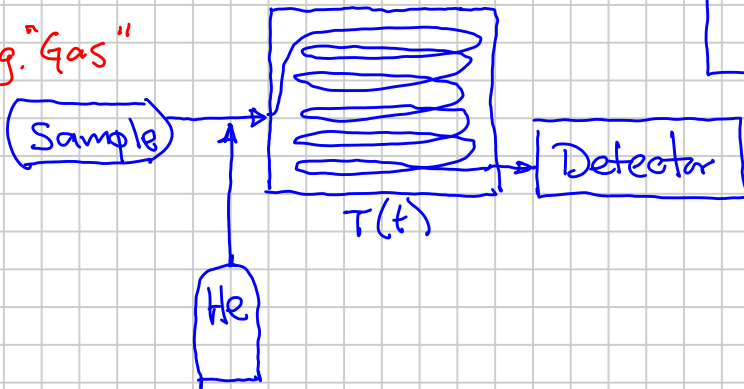
- Small Sample
- Helium carrier gas
- Packed Column
- Heater  $T(t)$
- Detector

Response

✓ (no uncertainty)



e.g. "Gas"



$$w_i = \frac{A_i}{A_T}$$

$M_i$  ✓ pure compounds

$$y_{gi} = \frac{w_i / M_i}{\sum_j w_j / M_j}$$

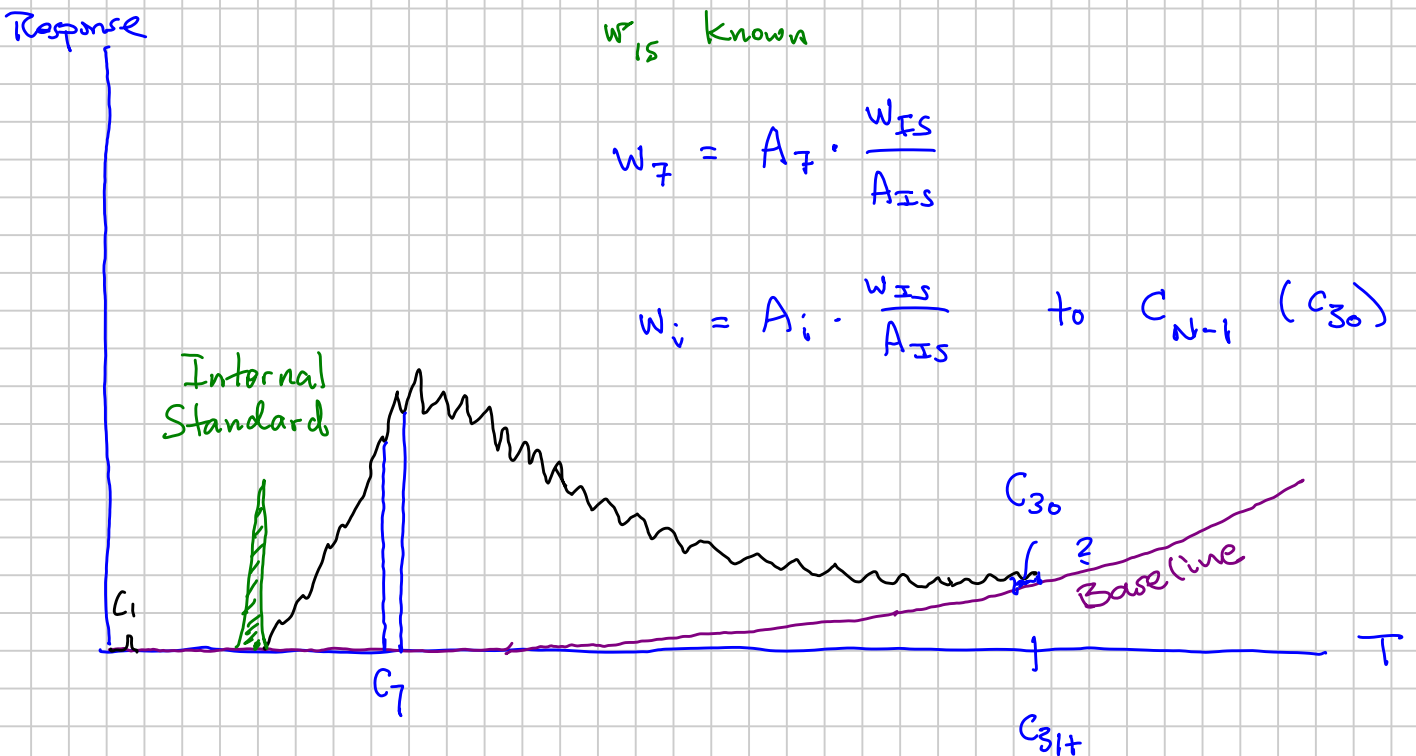
Flashed Oil Sample + I.S.

- He only

$w_{IS}$  known

$$w_7 = A_7 \cdot \frac{w_{IS}}{A_{IS}}$$

$$w_i = A_i \cdot \frac{w_{IS}}{A_{IS}} \quad \text{to } C_{N-1} (C_{30})$$



$$w_{31+} = 1 - \sum_{i=1}^{30} w_i$$

mass fractions  $\left\{ \begin{array}{l} w_i \text{ increasing uncertainty at higher SCNs} \\ w_{N+} \text{ max uncertainty as } N \text{ becomes big} \end{array} \right.$

↓  
Mole Fractions

$$\pm z_i = \frac{(w_i/M_i)}{\sum (w_j/M_j)}$$

Reported

Sample-specific uncertainty needs to be considered in EAS tuning process.

If  $z_i \propto w_i$  given then you can calc.  $M_i$  used by the lab

$$M_i = \bar{M} \cdot \frac{w_i}{z_i} \Rightarrow \bar{M} = M_{c_1} \cdot \frac{z_{c_1}}{w_{c_1}}$$

e.g. get  $\bar{M}$  knowing  $M_i, w_i, z_i$  for a pure compound  
 $M_{c_1} = 16.04$

Use  $(M_i)_{\text{lab}}$  to convert reported mole fractions to Lab "weight fractions" (measured), try to honor  $w_i$

$$\textcircled{1} \bar{M} = (16.04) (36.47/6.24) = 93.5$$

$$\textcircled{2} \text{Find Lab } M_{c_6} = (93.5) (3.97/4.33) = 85.7 < M_{n-c_6}$$

$$\text{Calc. } \bar{M}_{7+} = \frac{(\sum z_i M_i)}{(\sum z_i)} = \text{Cell M3}$$

mass / moles

SCN  
Table

Summarize Composition Uncertainties (in EOS modeling)

- ① Flash GOR used in recombination ± 5%
- ②  $w_i$   $i > C_{15}$  because of baseline shift
- ③  $w_{N+}$  because of  $\frac{1}{1 - \sum_{i=1}^N w_i}$  differencing like numbers

$\pm$   
 $\rightarrow 1$  as  $N$  grows

Gamma  
Molar  
Dist.  
Model
- ④  $M_i$  convert  $w_i \rightarrow z_i$

Exception of PhaseComp you can't quantify effect ①-④ ± on EOS tuning

\* ①-④ ±  $\Rightarrow$  mol% ± 2 for  $C_6$  &  $C_{7+}$  Limit as  $z_i$  change

\* Change in reported  $z_i$  should improve EOS predictions for ~ all PVT data ( $P_b, \rho, V_{no}, \dots$ )

PVTsim (others) to match  $p_s$  ( $p_d \neq p_b$ )

allow one molar composition regression variable:

$M_{n+}$  (last input component)

Internally calculates lab  $w_i$ ; all components (also  $C_{n+}$ )  
always honored

Recalculates  $z_i$  from  $(w_i)_{lab}$  and new  $M_{n+}$

$$z_i = \frac{w_i / M_i}{\left( \sum_{j=1}^{n-1} \frac{w_j}{M_j} \right) + \frac{w_{n+}}{M_{n+}}}$$

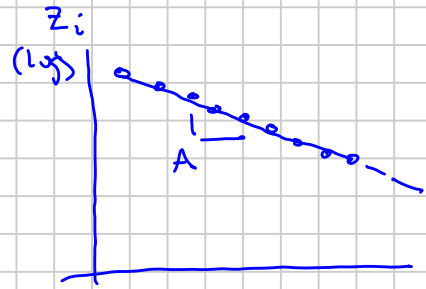
Every  $z_i$  changed



# Identify C<sub>7+</sub> Data Errors/Uncertainty

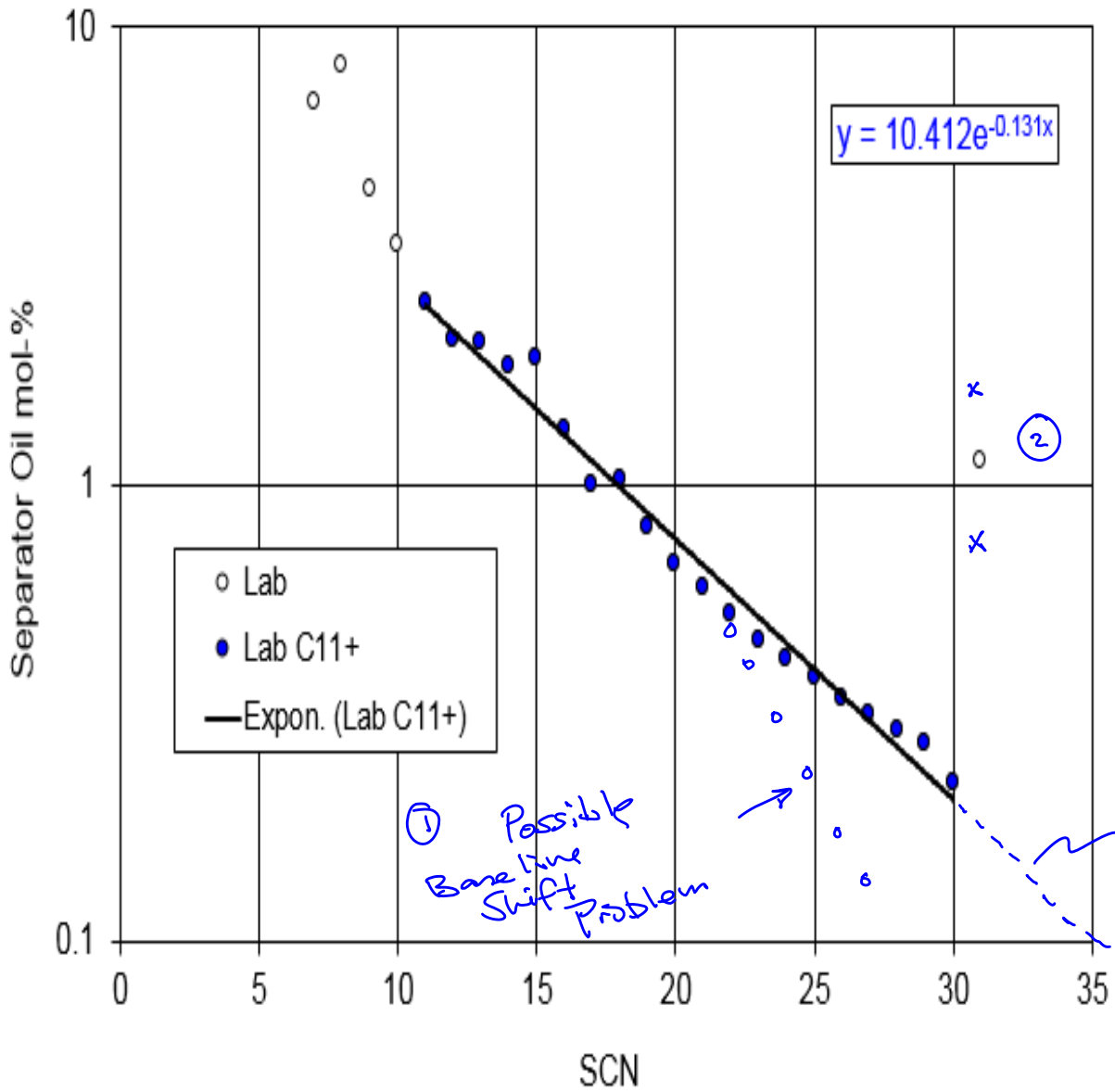
- Molar distribution model: Exponential

$$C_{n+} \quad Z_i = Z_n \cdot \exp [A(i-n)]$$



PVTsim

M:  
i



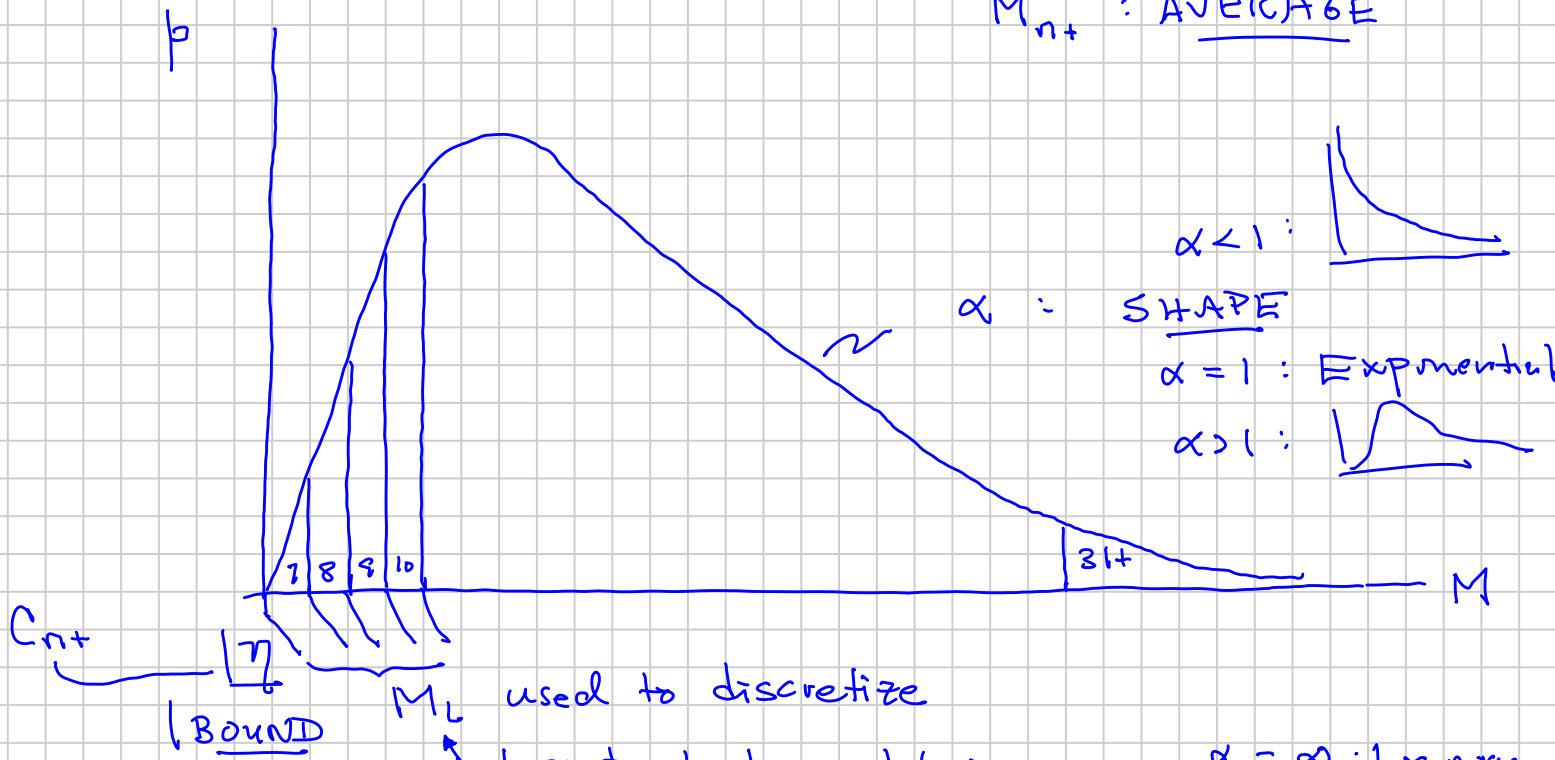
① Possible Baseline Shift Problem

Σ = x of C<sub>N+</sub>

$0.6 < \alpha < 2$  most reservoirs

- Gamma Distribution Model

$\bar{M}_{nt}$  : AVERAGE



$\alpha < 1$  :

$\alpha$  : SHAPE

$\alpha = 1$  : Exponential

$\alpha > 1$  :

used to discretize  
important to matching  
GC (TBP) data.

$\alpha = \infty$  : Log-norm.

$$\int_{?}^{\infty} p \, dM = 1$$

$$\hat{z}_i = \int_{M_{Li}}^{M_{Li+1}} p \, dM$$

$$m_i = \hat{z}_i \bar{M}_i$$

$$\bar{M}_i = \left( \int_{M_{Li}}^{M_{Li+1}} p \cdot M \, dM \right) / \hat{z}_i$$

mass / moles

$$\bar{M}_{nt} = \frac{\int_{?}^{\infty} p \cdot M \, dM}{\int_{?}^{\infty} p \, dM = 1}$$

$$\int_{?}^{\infty} p \, dM = 1$$

# Exponential (PVTsim) - discrete SCN model

$$M_i = 14 \cdot i + b$$

$$\text{PVTsim: } h = -4$$

$$\hat{z}_i = z_n \cdot \exp \left[ A(i-n) \right]$$

$\uparrow$                      $\uparrow$   
 $\bar{M}_{nt}$                 BOUND  
AVERAGE

$$\text{SHAPE} = 1$$

How to match GC (TBP) data to Gamma Model

① Only GC data

$w_i$   $z_i$  but not  $M_i$

Matching Amounts

\* Set  $M_{wi}$  fixed (how?)

Minimize mismatch between GC  $z_i$  ( $w_i$ )  
and the Gamma model values  $z_i$  ( $w_i$ ) } Find  $\gamma$ ,  $M$ ,  $\alpha$

② TBP data  $\underline{w_i}$   $z_i$   $(\bar{M}_i)_{\text{Lab}}$

Guess  $\Gamma$   $(\alpha, \gamma, \bar{M})$

$\uparrow$     $\uparrow$     $\uparrow$

minimize  $(\bar{M}_i)_{\text{Lab}}$  vs  $(\bar{M}_i)_{\Gamma}$

Find  $M_{wi}$  to get an exact match of  $w_i$  (or  $z_i$ )

$\Rightarrow (\bar{M}_i)_{\Gamma}$  from gamma model

Matching  $\bar{M}_i$  ( $Mw$ )

# SAMPLING

## \* Methods of Sampling

- ① 0 - Surface separator } Mobility-Average  
② 0 - Cased hole BHS } (Production test to Surface)  
                                  } Average over perf'd interval
- ③ 0 - Openhole Formation Testers (MDT) BHS @ Depth  
    - Preferred for Fluid Initialization  $z; (I, J, K)$
- { - Wellhead samples | Thornton | Petrotech } Unusual

## \* Why Sample?

① Collect samples for PVT measurements

to BUILD or VERIFY or IMPROVE an EOS Model

These samples do NOT need to be In-Situ Representative

~ Most all samples EXCEPT OBM-contaminated

② Compositions to help in Fluid Initialization

(In-Situ Fluid Mapping)

- All samples

- OFT samples are best

(Uncontaminated)

$z_s (I, J, K)$

$R_s, r_s (I, J, K)$

SPE 28829 : ECM

Gas Cap - Oil (Saturated)

# Separator Sampling in Gas Condensates (Vol. oils)

$GOR > 2000$  scf/STB

Recombination of Sep Gas & Sep Oil  $\Rightarrow$  Wellstream

should use

$$GOR_{sp} = \frac{q_{gspl} \text{ [scf/D]} \leftarrow}{q_{osp1} \text{ [sep-bbl/D sep-m}^3\text{/d]}}$$

Get this Right!

often report

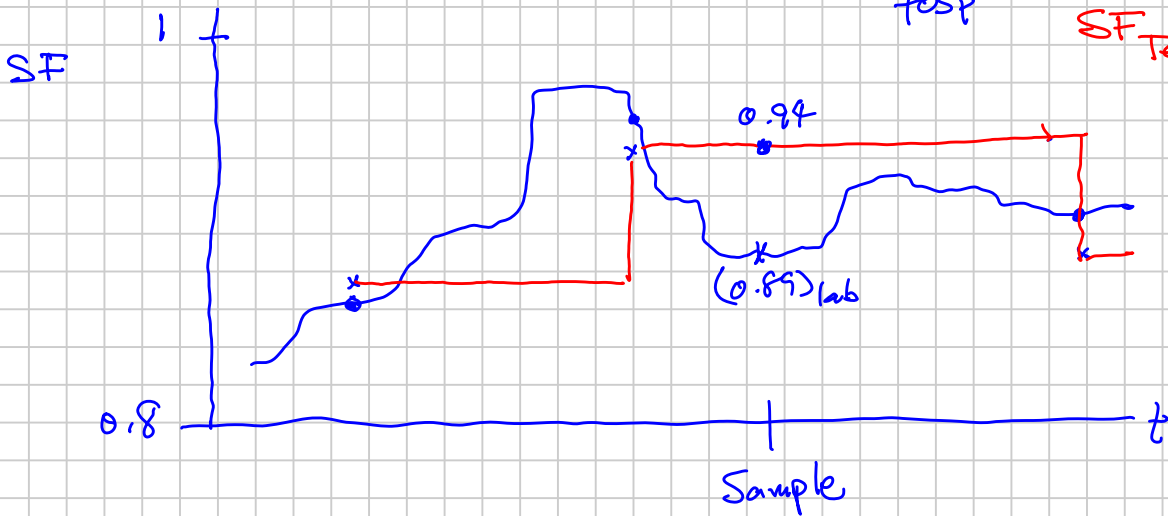
$q_o$  STB/D

$$q_o = q_{osp1} \cdot SF(\text{spot}) \quad \text{Test Company}$$

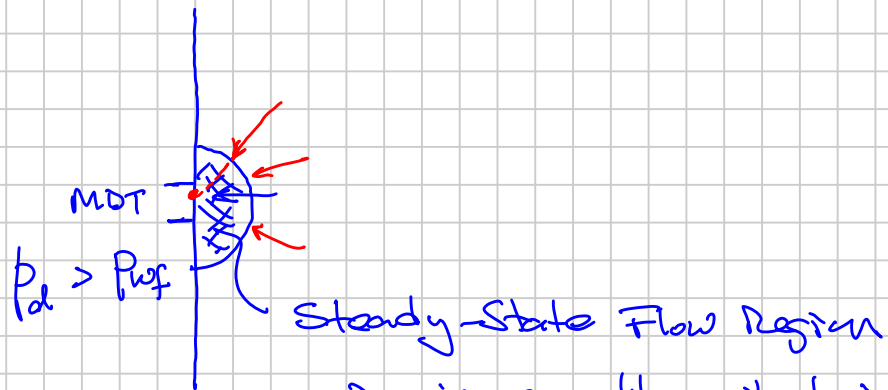
$$q_o = q_{osp1} \cdot SF(\underline{T}_{sp}, \underline{P}_{sp}) \quad \text{Correct}$$

metered

$$q_{osp} = \frac{(q_o)_{\text{Test}}}{SF_{\text{Test}}}$$



## Gas Condensate



$\lambda_0$  is exactly what is needed

to guarantee that what flows into the region, flows out of the region (into the MDT tool)