

### Free fatty acids in plasma:

all fatty acids (FAs) NOT bound in tryglycerids, phospholpids, cholesterol etc., hence "free": FFA

#### Is 'free' same as dissolved?

Practical insolvable: mainly bound in plasma albumin: FFA transporter

FFA-albumin → free albumin + dFFA

They should form the micelles and finally surfactant layer.

#### Problem:

Is FFA well (monomolecular) disolved?

# What is concentration of dFFA?

Some basic chemistry

reaction equation:

 $K_d = \frac{[free \ albumin] \cdot [dFFA]}{[FFA-albumin]}$ 

(K<sub>d</sub>: reaction constant of dissociation is 14 nM)

[free albumin] and [FFA total] can be measured,

Further:

Free albumin and FFA-albumin ca 106 x dFFA and K<sub>d</sub>,

So, [FFA-albumin] = [FFA] and [free albumin] = [total albumin] – [FFA].

$$[dFFA] = \frac{K_d \cdot [FFA]}{[total \ albumin] - [FFA]},$$

Very few is monomolecular dissolved, **dFFA** is in nM range!! (even less as bimeres etc.).

Detectable bubbles arise within ½ hour after surfacing. (long-chain) dFFAs should form monolayers to cover the bubbles.

#### **Problems:**

It there enough dFFA??
Are skin formed fast enough??

#### **QUESTIONS**

A What is the total bubble area to be covered?

B Is the quantity of dFFAs enough to cover all nuclei and bubbles?

C If not, is reservoir of FFA-albumin sufficient to cover all nuclei and bubbles?

D Is dissociation of FFA-albumin fast enough when amount of dFFAs is insufficient?

E Can the generation of all monolayers be completed in about one hour?

#### **QUESTIONS**

A What is the total bubble area to be covered?

Suppose: Bubble grade is KM = 1:

detectable bubbles ca. 2 /L (surface  $10^{-1}$  mm<sup>2</sup>, D = 180  $\mu$ m),

micro-bubbles about 10<sup>5</sup> with a surface of 10<sup>-3</sup> mm<sup>2</sup>,

about 10<sup>7</sup> with 10<sup>-5</sup> mm<sup>2</sup>,

Nuclei about 10<sup>9</sup> with 10<sup>-7</sup> mm<sup>2</sup>?

Together: 300 mm<sup>2</sup>/L plasma

Bubble grade is KM = 4: Together: 6000 mm<sup>2</sup>/L?

## **QUESTIONS**

A What is the total bubble area to be covered?

B Is the quantity of dFFAs enough to cover all nuclei and bubbles?

With some 4 nm/L (outcome reaction equation) ca. 275 **mm²/L plasma** 

Even for KM = 1 this is hardly/not enough.

#### **QUESTIONS**

A What is the total bubble area to be covered?

B Is the quantity of dFFAs enough to cover all nuclei and bubbles?

C If not, is reservoir of FFA-albumin sufficient to cover all nuclei and bubbles?

FFA-albumin is in the mM range, so

for KM = 1 excess is factor of some 40.000 X

for KM = 4 some 1000x

#### **QUESTIONS**

- A What is the total bubble area to be covered?
- B Is the quantity of dFFAs enough to cover all nuclei and bubbles?
- C If not, is reservoir of FFA-albumin sufficient to cover all nuclei and bubbles?
- D Is dissociation of FFA-albumin fast enough when amount of dFFAs is insufficient?

Dissociation of FFA-albumin is often needed.
Rate of dissociation is can be 1.4 mmole/s: immediate replenishment since micelle formation is very slow.
So, dFFA is always the same for given total [albumin[ and [FFA].

#### **QUESTIONS**

A What is the total bubble area to be covered?

B Is the quantity of dFFAs enough to cover all nuclei and bubbles?

C If not, is reservoir of FFA-albumin sufficient to cover all nuclei and bubbles?

D Is dissociation of FFA-albumin fast enough when amount of dFFAs is insufficient?

E Can the generation of all monolayers be completed in about one hour?

No, takes possibly hours to combine mono-, bi- etc polymeres of FFA to micelles and then to complete skins. The **CMC (critical mycel concentration)** to form mycelles is about 1 mM of dFFA, close to a million times the actual dFFA concentration!

# **Conclusion from theoretical study**

**Long-chain** FFA can not be used to form surfactant skins around DCI bubbles.

**Medium-chain** FFAs are better dissolvable and their CMC are much higher. However, **they do not occur in food** or the concentration is much lower than CMC (Naoctanoic acid 0.36 M).

Micelles of **short-chain** FFAs do occur and with n=4 or 6 CMC is very high and it is highly questionable whether they can form stable skins.



# Methods



52 male divers

precordial Doppler method, 40, 80, 120 and 160 min) → Kisman Integrated Severity Score (logKISS).

Half of subjects obtained fat rich **and** half fat poor meals to enlarge the FFA and TriG range of blood plasma.

63 simulated dives (21msw/40min profile)

11 both (paired testing).

Correlate post exposure, dFFAs and total FFA (mM range), with **venous gas bubbles** (KISS at 40, 80, 120, 160 min post-dive, precordial)

- Physics

am	Methods								
	Gro	up Frich, n=	=28	Gro	oup Fpoor, n=	=24			
	Age	VO <sub>2max</sub>	body fat	Age	VO <sub>2max</sub>	body fat			
	(years)	(ml/kg.min)	(%)	(years)	(ml/kg.min)	(%)			
mean	45.5	42.3	21.2	46.3	42.0	21.2			
SD	3.47	4.82	2.9	3.10	6.62	4.2			

Statistics: groups perfectly matched

Δ's: 0.8 year (2%), 0.3 ml/kg.min (1%), 0.0% BF

DOM PROM



# Group differences post exposure



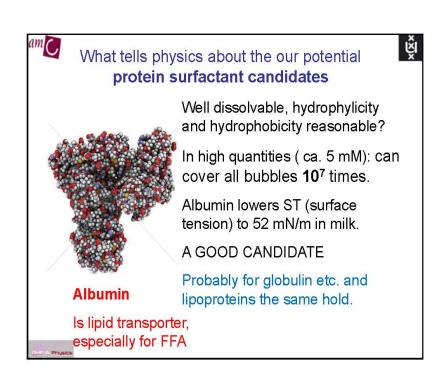
Measured item	Fat-rich group, n=28	Fat-poor group, n=24	Frich - Fpoor	
	mean± SD	mean±SD	p-value t test	
FFA	0.20±0.08	0.078±0.04 9	7x10-8	

# Results

No correlation between post exposure albumin, dFFAs (nM range) and total FFA (mM range) and **bubbles.** 

Also not with TriG and TCh

33







# MEASURING SURFACE TENSION

When a substance lowers surface tension  $\rightarrow$  decreases R<sup>crit</sup>, hence bubbles have longer time to grow  $\rightarrow$ 

more bubble survival  $\rightarrow$  more bubbles.

DATE OF PRODUCTS



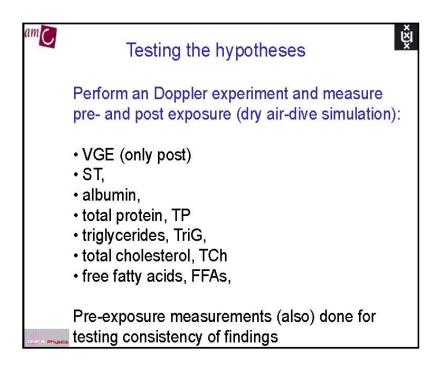


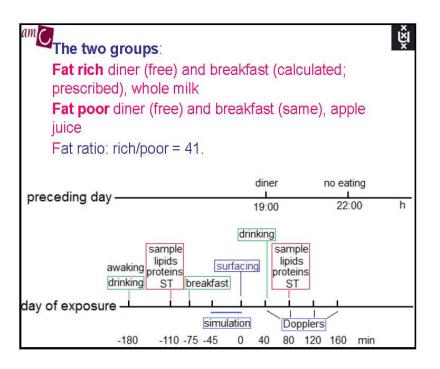
# MEASURING SURFACE TENSION

# The hypotheses

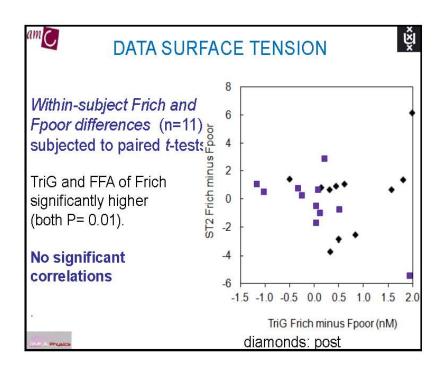
We expect the following associations, i.e. with significant correlations:

- Lipids&proteins (FFAs etc. and albumin) with ST (negative correlate).
- Proteins (albumin) with VGE (negative correlate).
- ST with VGE (negative correlate).





ndependent -> dependent ↓	TCh	TGI	FFA	alb	TPr	ST
ST	079 <b>.55</b>	0.078 .56	16 .24*	23 .09	.31 . <b>02</b> **	
logKISS	17 .20	.17 .20	07 .60	.01 .92	.02 .88	.01 .93
** not signification Only Total exposure Bubbles substan there are	Prote no sig are i ce or	in seer nificand not aft by S	ns to a ce. He fecter [:	affect nce r d by	ST bu	





# **Main Findings**



- No significant and consistent effects of lipids and proteins on ST pre- and post exposure.
- · Lipids and proteins do not affect VGE.
- VGE does not correlate with ST.
   (see also Gempp et al, Br J Sports Med 2009)
- All analyses with subjects with KISS>0: same results
- y is ca. 57 mN/m (corrected).
- No KISS differences found with within-subject fatrich versus fat-poor meals (paired t-tests, no significant correlations).

C. C. Physics

# am C

# **Discussion 1**



FFA's etc. are not good surfactant candidates,

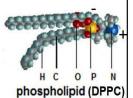
But what about phospholipids?

All have very poor solubility and high CMCs,

Except DPPC (dipalmitoylphosphatidylcholine):

hardly soluble and low CMC.

Can it form dimeres etc. in plasma?



Anyway: for DPPC-skin ST is much too high

real Physic





### Discussion 2

ST higher than expected, ca. 57 mN/m (corrected)

→ small stabilizing effect (r<sup>crit</sup> ca. 20% lower) too small for effect on KM

Yount 20 mN/m: 2½ x more bubbles! Well measurable.

Possibly, the 15 mN/m decrease (rel. to water) is caused by predominantly protein mixture, surrounding the bubbles.

Albumin and tot-protein levels practically invariable (post-pre, rich-poor & within-subject)!!

Division



# **Discussion 3**



Albumin is a promising candidate (milk chemistry) to coat bubbles and such reducing **y**.

It has 9 binding sites for FFA and it also bounds phospholipids. Their C-tail may point to the bubble interface.

DPPC is probably also embedded in albumin. But indissoluble DPPC multimeres and micelles from membrane destruciion can be suspended in the plasma.

PTUN

