

Transport Phenomena.
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Lecture-43.
Mass Transfer.

So far we have discussed momentum transfer and heat transfer and we have also worked what would be the type of transport, how can we press mathematically the transport inside a boundary layer, a hydrodynamic boundary layer and inside it a thermal boundary layer. For the case of hydrodynamic boundary layer we have seen how to obtain an analytical solution by converting a PDE which was originated out of the simplification of Navier Stokes equation for flow inside a boundary layer on a flat plate.

So we used the method of combination of variables to obtain not only the velocity V_X and V_Y inside the boundary layer, but also the gradient of velocity or at Y equals 0, that means that liquid solid interface, which was then used to obtain an expression for the friction coefficient C_f . We then proceeded for solution of the or treatment of the turbulent boundary layers. And in turbulent boundary layers we saw that the situation became so complicated that it is not possible even to write the governing equation, the statistically it is having an universal velocity profile which would be valid in all regions of turbulent flow for flow over, even for a flow over a flat plate is extremely difficult.

So we have seen dividing the flow regime into 3 layers like viscous sublayer, the transition region and a turbulent core for the case of pipe flow. For each of these regions, there was a velocity profile. And there was also an entirely empirical velocity profile known as the one 7th power law which can be used to fit the experimental data, especially for those points situated near the centre. However this expression, even though it fits the data wells, it fails on the solid wall because you cannot evaluate the gradient of the velocity at Y equals 0, that means on the flat plate using one 7th power law, that was a major limitation.

So then we proceeded to obtain a momentum integral approach, an integral approach which resulted in a momentum integral equation and there with some approximation with an assumed value, assumed expression of velocity profile where the constants were evaluated with the boundary conditions that we have on the plate, namely no slip condition and at the edge of the boundary layer, that means where the velocity in the X direction would be equal to the free stream velocity and the velocity profile approaches the free stream with 0 slope. Using these conditions and assumed velocity profile, we could move ahead to obtain the,

what would be the boundary layer growth in the case of turbulent flow and what would be the friction factor expression.

We then used the same concepts for the treatment of the thermal boundary layer. In thermal boundary layer before that it is the equation of energy applicable inside the thermal boundary layer after the standard simplifications which are boundary layer approximations, this equation is coupled to the velocity boundary layer equation, namely the Navier Stokes equation for flow inside the boundary layer. The coupling appears because of the appearance of velocity V_X and V_Y in the energy equation. So it is a one-way coupling.

We showed in previous classes how to solve this thermal boundary layer growth and the velocity, when the temperature gradient at the solid liquid interface through the simultaneous solution of the momentum equation as well as the energy equation. So for laminar flow we have obtained an expression for Nusselt number which is the convective heat transfer coefficient multiplied by length scale divided by the thermal conductivity of the fluid, this Nusselt number was related to 2 dimensionless groups which appeared automatically through the nondimensionalization of the governing equation, namely Reynolds number and the Prandtl number.

So we got compact expressions of Nusselt number, the, with the Engineering parameter convective heat transfer coefficient embedded into the Nusselt number. So that is, the H is the one that we would like to evaluate and the corresponding dimensionless group is Nusselt number. So we got relationship between Nusselt number, Prandtl number and Reynolds number with a constant in front of it. But that was for laminar flow, I did not say anything about the turbulent flow inside a thermal boundary layer.

The treatment of turbulent flow inside a thermal boundary layer is more complicated, more complex because you are going to have transfer of heat not only by conducting and convection, there will be the formation of eddies and these eddies. And these eddies would carry additional heat, additional energy for flow when for the case when the flow inside the boundary layer turns to be turbulent. So the presence of eddies creates or imposes additional problems in solving the energy equation. 1st of all we do not know what would be the right form of energy equation.

So even if we express the energy equation in the same way as we have done for the case of momentum boundary layer, that is in terms of fluctuating components, it is almost intractable.

So we have to think of some ways to use the solutions that we have already obtained for the case of momentum boundary layer, both in laminar flow as well as in turbulent flow is there any way to use, to project those relations and or correlations for the case of turbulent flow in thermal boundary layer. So that means I am trying to find an analogy, a logical set of conditions which must be met, so that the results of momentum transfer in turbulent flow can be projected and to obtain the results connecting the dimensionless groups, relevant dimensionless groups for the case of turbulent flow inside a thermal boundary layer.

And if we can establish this transformation, then the same logic can also be obtained, can also be used to obtain the relation between the relevant parameters in for mass transfer inside the concentration boundary layer. The way we have the velocity boundary layer we have seen what is the thermal boundary layer. Similar to thermal boundary layer, I will also have the concentration boundary layer in which the species concentration would change from some value on the solid plate to a constant value in the free stream of the flowing solution above the solid plate.

So the results of, so it would be, it would be the objectives, as I mentioned at the beginning of the course the objective of this course would be how, why and when we can transform the relations obtained in hydrodynamic part of the boundary, solution of the hydrodynamic boundary layer, how can you use that as a solution of the thermal boundary layer and then for the concentration or the mass transfer boundary layer. So I am not going to do that right now, that would be the last topic of this course. So I would very quickly go through some of the salient features of the mass transfer process which is complicated because now we are dealing with mixtures of at least 2 components, maybe a solute and a solvent.

So at least 2 species are present in the case when we are having net transport of one species from the one point to the other. Now the net transport asks the species from one point to the other exactly like in the case of heat transfer, it can take place because of actual flow from point A to point B which carries component 1 from A to B. So that is due to the imposed flow of the solutions from point A to point B carrying component A from one location to the other, which is nothing but the convective motion of the species, the motion of the species due to convection imposed on the flow field.

There would be another way by which mass gets transported which is similar to the conduction heat transfer, heat transfer by conduction. So whenever there is a temperature gradient, even if there is no flow we will still have heat transfer because of molecular,

because of mobile means, because of molecular mechanism. So this conductive heat transfer which depends not only on the concentration difference but on the gradient of concentration between, gradient of temperature between 2 points, exactly similar phenomena exists for mass transfer as well which are aptly called the diffusive mass transfer.

So in diffusion or diffusive mass transfer, mass travel from one location to the other if there is an imposed concentration gradient, the concentration gradient may exist as a result of several conditions. But if there is a temperature, if there is a concentration gradient, then mass gets transferred from one point to the other. So similar to heat flux, similar to Fourier's law, similar to Newton's law of viscosity, the mass flux as in the case of heat flux is proportional to the concentration gradient. Think of this similarity with the temperature gradient.

So mass J is proportional to the concentration gradient and the proportionality constant with a - sign, since mass always travels from high concentration to low concentration, the concept of this expression is commonly known as the diffusivity of one in two. It is expressed in D , in the form D with a subscript AB which is the constant, which is the diffusion coefficient of A in B. Now some of the relations this is known as the Fick's law of diffusion which like Newton's law or Fourier's law is a phenomenological equation, it cannot be derived, it can, it was arrived by looking at the data of many experiments over a large range of concentrations and it was found that a mass flux is always proportional to the concentration gradient.

So before we use Fick's law and other physical boundary conditions in solving, in modelling the mass transfer process, the same way we have done for heat transfer, it would be, I would like to go quickly through the relations, the established relations in mass transfer that I am sure you already know, it is only going to be a recapitulation of what you have studied in your mass transfer 1 or mass transfer 2 courses. So our study of mass transfer and modelling of processes involving mass transfer, the mass transport process, we understand, we start with the fact, with the with the realisation that mass transfer is a more complicated process as compared to a momentum transfer all heat transfer because more than one species is involved.

So you will have at least 2 species and in multicomponent systems, we will have more than one species, more than 2 species which are present in the system and therefore the, not only the diffusion coefficients are going to be different, however the motion of the one species, the motion of the molecules of one species in a medium will start to affect the molecules of the

and we understand that X_A which is a mole fraction, $X_A + X_B$ equal to 1, the same way weight fraction sum would be equal to 1 and these relations are self-explanatory.

So we have all seen these expressions before and we will see how these expressions will be used later on to understand, to express the mass transport process in a system where we have both convection as well as the diffusive mass transfer. So coming back to some of the other equations, other definitions that one can think of, one uses in mass transfer process.

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LOCAL MASS AV. VEL $v = \frac{\sum_{i=1}^n \rho_i v_i}{\sum_{i=1}^n \rho_i}$

LOCAL MOLAR AV. VEL $v^* = \frac{\sum_{i=1}^n C_i v_i}{\sum_{i=1}^n C_i}$

DIFFUSION VELOCITIES

$v_i - v = \text{DIFF}^n \text{VEL. of } i \text{ w.r.t. } v$

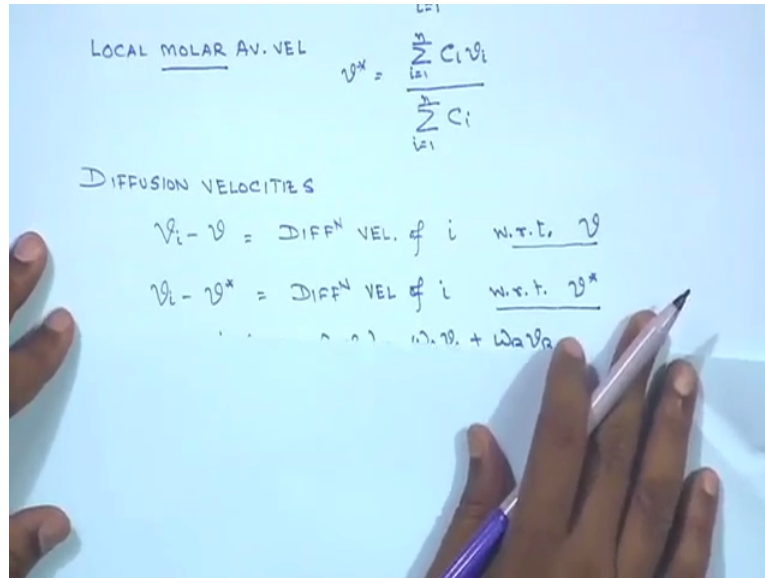
$v_i - v^* = \text{DIFF}^n \text{VEL. of } i \text{ w.r.t. } v^*$

There is something called local mass average velocity which is simply $\rho_i v_i$ divided by summation of ρ_i overall the species. So this is the mass average velocity, the same way you have the mass average velocity, you can also find out, will also express the molar average velocity where simply the mass concentration is replaced by the molar concentration. So the denominator is the total mass concentration whereas the denominator over here is the total molar concentration of the solution. Now whenever something, whenever a component moves in a, moves in the solution, then you can either fix the coordinate systems and keep them stationary.

So the 2 definitions of mass average velocity and molar average velocity that I have shown you before are with respect to stationary axes. Now the mass average velocity can also be used, let us say I have a portion of a solution in which there is a diffusing species A and this entire species has some mass average velocity with which it is let us say moving in this direction. However the species A present in it has a different velocity because it is also

diffusing as a result of the bulk flow as well as as a result of a concentration gradient imposed on it.

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So if we want to separate the diffusion from the bulk motion of the fluid, then the diffusion velocity is something which is exclusively due to diffusion for a species A and therefore the diffusion velocity is expressed as V_i , that is the velocity of the i th species, subtract from that the local mass average velocity, so the difference in velocity, the additional velocity that the i th component has is over the mass average velocity is termed as the diffusion velocity. So V_i , the velocity of the i th species - the local mass average velocity is termed as the diffusion velocity of i with respect to V .

Now the way you have expressed the diffusion velocity where the basis is taken as the local mass average velocity, you can take the basis as local molar average as in V^* . So the component velocity, the relative velocity of the component can be expressed with respect to the mass average velocity or with respect to the molar average velocity, both are diffusion velocities, one with respect to V , the other is with respect to the molar average velocity. So these are the 2 diffusion velocities that are commonly used for, in case of, in for mass transfer.

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$$v^* = \frac{\sum_{i=1}^n c_i v_i}{\sum_{i=1}^n c_i}$$

DIFFUSION VELOCITIES

$$v_i - v = \text{DIFF}^n \text{ VEL. of } i \text{ w.r.t. } v$$

$$v_i - v^* = \text{DIFF}^n \text{ VEL of } i \text{ w.r.t. } v^*$$

$$v = \frac{1}{\rho} (\rho_A v_A + \rho_B v_B) = \omega_A v_A + \omega_B v_B$$

$$v^* = \frac{1}{c} (c_A v_A + c_B v_B) = x_A v_A + x_B v_B$$

And this velocity, the velocity of this if you expand it, is simply going to be 1 by rho, 1 by rho times rho A VA + rho B VB in which if you take the ratio of rho A by rho, it is simply going to be the weight the weight fraction of component A and similarly V Star which is a local molar average velocity. If you expand this, it is simply going to be 1 by total concentration, then molar concentration of A, molar concentration of B and the velocities of A and B and CA by C is nothing but the fraction, the mole fraction of component A and mole fraction of component B. So these 2 relations directly follow from the definition of the mass average velocity or the molar average velocity.

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MASS & MOLAR FLUXES REL. TO STATIONARY COORDINATE S

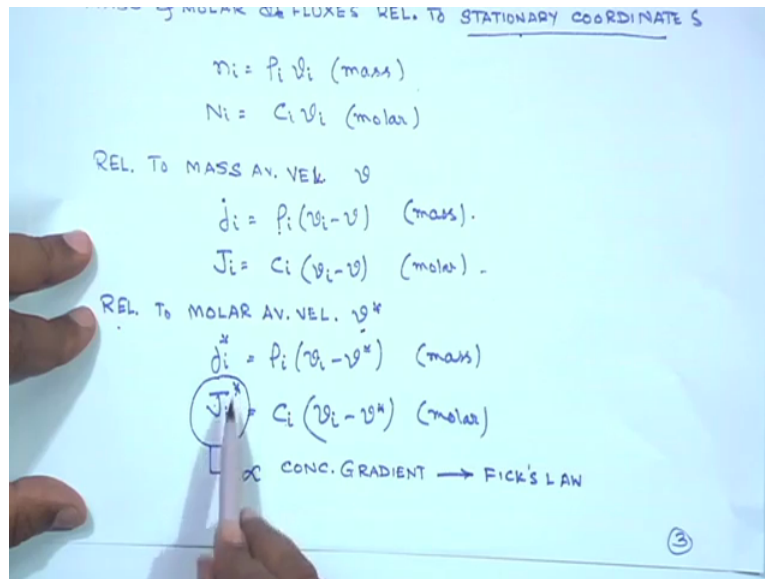
$$n_i = \rho_i v_i \text{ (mass)}$$

$$N_i = c_i v_i \text{ (molar)}$$

REL. TO MASS AV. VEL. v

$$j_i = \rho_i (v_i - v) \text{ (mass)}$$

$$J_i = c_i (v_i - v) \text{ (molar)}$$



Now since we have defined the diffusion velocities in this way, the diffusion velocities can then be converted to fluxes. So one would be a molar flux and the other would be mass +, so in terms of stationary coordinates, it is simply going to be $\rho_i v_i$, if you express it in terms of mass or if express in terms of moles, it is simply going to be $C_i v_i$, these v_i s are with respect to stationary coordinates. When you bring the same mass flux or molar flux and take the average velocity, average velocity \bar{v} to be the basis, so relative to the mass average velocity, the mass flux or the molar flux denoted by j or capital J for the i th species is simply going to be ρ_i times the relative velocity square for the relative velocity we use the mass average velocity.

So it can either be expressed in terms of mass or it can be expressed in terms of moles. Same thing if we do in terms of molar average velocity, so this is in terms of stationary coordinates where the velocity is 0, where \bar{v} is 0, so it is $\rho_i v_i$. If I have a mass average velocity and I express relative to the mass average velocity, it is simply going to be $v_i - \bar{v}$. If I do it in terms of molar average velocity, it is going to be $v_i - \bar{v}^*$ for the case of mass and for the case of moles it is simply going to be C_i .

Now this flux, be it mass flux or molar flux is generally expressed, the molar flux where J_i^* if you look at J_i^* , this is the molar flux of the i th species when the flux is relative to the molar average velocity which is, so $J_i^* = C_i (v_i - \bar{v}^*)$, this molar flux is proportional to the concentration gradient. That is the statement of Fick's law. So Fick's law essentially tells us that the molar flux J_i^* is proportional to the concentration gradient. So Fick's law is simply the flux, molar flux of component A when expressed with respect to the average, molar average velocity is simply $-D_{AB}$ times divergence of X_A .

So if C is constant, I can put C inside, it is simply the other way of writing it is time CA . So this for a rectangular coordinate system can be written as $-DAB \text{ times } \text{Dell } CA \text{ Dell } X + \text{Dell } CA \text{ Dell } Y + \text{Dell } CA \text{ Dell } Z$. So see the similarity that we have for the case, for decay with heat transfer and with mass transfer. So this is the statement of Fick's law and from our definition of molar flux, this is the definition of molar flux when expressed in terms of V star. So I expand it and $CA VA$ is nothing but the smaller flux of A and then when you expressed this V Star, the formula would simply be $CAVA + CB VB$ by C . CA by C is the molar, the mole fraction, so your JA is equal to $NA - XA \text{ times } NA + NB$.

Plug it in here and what you have is another form of, another form of Fick's law. So this is, this NA is relative to stationary coordinate and what it tells is that the molar flux of component A relative to stationary coordinates from here is a sum, is an algebraic sum of $XA \text{ times } NA + NB$ where $XA \text{ times } NA + NB$ is the flux due to the bulk motion of the fluid. So $CA \text{ Times } V \text{ star}$, $V \text{ Star}$ denotes the molar average velocity. So if we have a bulk velocity present, bulk motion present in the fluid, it is also going to contribute to a flux of A .

As I said the species A can move from point A to point B if there is a bulk motion. There may not be any difference in concentration, so a sugar solution, a constant concentration sugar solution may be forced, may be allowed to move from point A to point B by imposing a pressure gradient. There is no diffusion, since the concentration is same everywhere, but what you have a bulk motion of the sugar molecules from point A to point B . So this kind of bulk motion imposed by the flow only is the significance of the 1st that we have here which is $XA \text{ times } NA + NB$.

So this is due to bulk motion. Sometimes in addition to bulk motion or even in the absence of bulk motion you have concentration difference. So if you have concentration difference or more correctly if you have a concentration gradient present in the system, then this is going to give rise to say diffusive motion of A , species A . So the total effective motion of species A is the algebraic sum of the species movement due to bulk motion and or the species movement due to the concentration gradient imposed due to certain conditions present in it. So therefore the problem of Fick's law is to be resolved, the 1st thing that needs to be resolved is what is, how to get rid of NB from the expression of NA . So NA is $XA NA + NB - DAB \text{ Dell of concentration of } A$.

So 1st of all it may be mentioned that it is diffusion only process, that means there is no imposed bulk flow. So if it is a diffusion only process, then the 1st term on the right-hand side

which signifies bulk motion can be dropped. So that is one way of getting rid of N_B which is the unknown, which is which appears in this expression. So if it is a bulk motion, if it is a diffusive motion only situation, then this can be dropped. In some cases there would be a relation between N_A and N_B which arises due to some other factors.

For example it could be case of equimolar counter diffusion. That means for one mole of A moving in this direction, one mole of B is moving in the opposite direction. So if this is a case of equimolar counter diffusion where N_A is going to be equal to $-N_B$, for that specific case N_A is going to be equal to $-N_B$ and therefore this term, the contribution of this term would be 0. So the expression would be same as that of diffusive motion only situation but for different reasons. Since it is equimolar counter diffusion, N_A and N_B would cancel out each other.

There are, in some cases the stoichiometry of the reaction if it is a reacting system, let us say 3 moles of A comes and reacts on a catalyst surface generating 2 moles of B which then travel in the reverse direction towards the bulk. So for every 3 molecules of A coming to a specific direction 2 molecules of B would have to travel in the opposite direction at steady-state in order to maintain the concentration at each point, either independent of time. So the concentration of A may vary, concentration of B may vary but the concentration at the fixed location is not, will not vary with time. So that is that is what the steady-state is.

So in some cases stoichiometry of the reaction, stoichiometry of the reaction taking place between 2 reacting components would give you some idea between the relation between what, how N_A is related to N_B . So in absence of any such generalisation, any such simplification, the expression to be used for the molar flux of component A will consist of 2 terms, one due to bulk motion and other due to the concentration gradient. So this D_{AB} which is the diffusion coefficient of A and B for, they behave slightly differently for gases and for liquids, okay.

So most of them increase with an increase in concentration in temperature, so as the temperature increases, this D_{AB} , it is a function of pressure, temperature and it is also a function of, it could be function of composition, composition of the gas mixture. So the gases, further gases and liquids, with increasing temperature D_{AB} increases and at low-density it is almost composition independent for the case of or for the case of gases. So what we have then we need to see the similarity between D_{AB} which is the mass diffusion

coefficient expressed either in terms of mass or in terms of mole, compare that with γ which is μ by ρ and compare that with K by ρC_p which is α .

So if you compare D_{AB} , the diffusion coefficient of A and B, μ by ρ , which is the kinematic viscosity and K by ρC_p which is the thermal diffusivity, so momentum diffusivity, thermal diffusivity and mass diffusivity, all will have units of metre square for second. So these 3 are similar in nature, the mass diffusivity, the momentum diffusivity and the thermal diffusivity, they have the same unit as metre square per second and all of them denote the transport of mass, the transport of momentum or the transport of energy when you impose a concentration gradient, velocity gradient or a temperature gradient.

So this would be the beginning, the start of the, start of the analogy, finding the analogy between different processes heat transfer, mass transfer and momentum transfer. But before we reach that point what I would do in the subsequent process, 3 or 4, 4 or 5 classes after today is to show you examples by which 1st of all a shell component balance can be used, can be used to obtain the concentration profile of a specific component in a system where it is a diffusion only process, where both diffusion and diffusion and bulk motion convection is present and different ways by which N_A can be related to N_B .

The molar flux of A and the molar flux of B, what is the relation between them apart from counter diffusion, equimolar counter diffusion and so on. And finally the, like heat generation in the case of energy question, we can also have generation of a species due to reaction, generation or depletion of species due to reaction in a medium in which A is diffusing. So if A is diffusing and as it diffuses, it reacts with another reactant B present in δ , then A is going to get depleted as it moves in the solution.

So reaction, chemical reaction, homogeneous chemical reaction can act as a source or a sink term in the mass balance equation. So when we write the shell balance of shell component balance, the same way we have done for, for the previous cases, heat transfer and momentum transfer. The source or sink term, for example in nuclear heat source or an electrical heat source, the equivalent of that in the case of mass transfer would be, if there is a reaction which is consuming A or a reaction which is producing A in the entire domain of transfer of A from point 1 to point 2.

So a homogeneous reaction would appear as a source or sink term in the governing equation. So contrary to that if it is a heterogeneous reaction, that means if it is a, let us say it is a

catalytic reactions well this is the catalyst surface and the reaction of A getting converted to B is going to take place only on the catalyst surface, it is a heterogeneous reaction. Therefore A diffuses and reaches at this point when it gets converted to B and the products will diffuse back to the mainstream. In that case, since in the path of diffusing A, it does not encounter any reaction, generation or depletion.

The generation or depletion takes place only at a specified location, that is on the catalyst, heterogeneous reaction, this condition would appear as a boundary condition in the governing equation. So we have to keep in mind the difference between the heterogeneous reaction and the homogeneous reaction, one in which it appears as a source or sink term in the governing equation itself and the other variant appears as a boundary condition. So we will see examples of that, examples of modelling the process in the coming 4 or 5 classes and then we will finally move to the final part of this course which is to see, to evaluate the analogy between heat, mass and momentum transfer.